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Minimum DB
Maximum DB
                                                                                                                                                                                                                                                                                                              Result
No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              on:
                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                              Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        seq length: 0
seq length: 2000000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Published Applications AA:*

1: /cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

3: /cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

4: /cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

6: /cgn2_6/ptodata/2/pubpaa/US08_NEW_PUB.pep:*

6: /cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*

7: /cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*

9: /cgn2_6/ptodata/2/pubpaa/US09A_PUBCOMB.pep:*

10: /cgn2_6/ptodata/2/pubpaa/US09A_PUBCOMB.pep:*

11: /cgn2_6/ptodata/2/pubpaa/US09A_PUBCOMB.pep:*

12: /cgn2_6/ptodata/2/pubpaa/US09A_PUBCOMB.pep:*

13: /cgn2_6/ptodata/2/pubpaa/US09A_PUBCOMB.pep:*

14: /cgn2_6/ptodata/2/pubpaa/US09A_PUBCOMB.pep:*

15: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

16: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

17: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

18: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

19: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

19: /cgn2_6/ptodata/2/pubpaa/US10B_PUBCOMB.pep:*

19: /cgn2_6/ptodata/2/pubpaa/US11A_PUBCOMB.pep:*

20: /cgn2_6/ptodata/2/pubpaa/US11A_PUB.pep:*

20: /cgn2_6/ptodata/2/pubpaa/US11A_PUB.pep:*

21: /cgn2_6/ptodata/2/pubpaa/US10B_PUB.pep:*

22: /cgn2_6/ptodata/2/pubpaa/US10B_PUB.pep:*
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Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                           Query
Match Length
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586
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1854112 seqs, 416015017 residues
            99.3
92.0
92.0
91.8
91.8
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91.8
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Copyright (c) 1993 - 2005 Compugen Ltd
       US-10-010-729-9
US-10-194-801C-4
US-10-719-461-50
US-10-322-673-53
US-10-981-621-53
US-10-981-673-53
US-10-981-673-53
US-10-981-691-53
US-10-727-155-310
US-0727-155-311
US-09-880-748-1260
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                                                                                                                                                                                                                                                                                                        Description
Sequence 9, Appli
Sequence 4, Appli
Sequence 50, Appl
Sequence 53, Appl
Sequence 310, Appl
Sequence 311, App
Sequence 1160, Ap
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ALIGNMENTS

	45	44	43	42	41	40	9	38	37	36	35	ω 4	ü	32	<u>ω</u>	30	29	28	27	26	25	24	23	22	21	20	19	18	17	16	15	14	μ	12
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	110	255	255	254	254	248	248	130	123	256	256	251	251	248	248	248	244	251	253	253	111	251	251	248	248	248	110	110	255	255	112	248	248	258
	15	15	10	15	10	20	16	18	14	15	10	15	10	18	15	10	16	17	15	10	14	15	10	16	15	10	17	17	15	10	14	15	10	15
	-10-440-522-7	-10-293-418-	-09-880-748-	-10-293-418-113	-09-880-748-	-11-090-84	-10-779-461-	-10-909-851-	-10-153-43	-10-293-418-	-09-880-748-	-10-293-418-	-09-880-748-	-10-981-692-	-10-293-418-	-09-880-748-146	-10-779-461-	-10-935-290-	-10-293-41	-09-880-748-185	-10-269-805-	-10-293-418-	-09-880-748-155	-10-779-461-21	-10-293-418-86	-09-880-748-86	-10-727-155-	-10-727-155-	-10-293-418-11	-09-880-748-	-10-269-805-10	-10-293-418-12	-09-880-748-124	US-10-293-418-1260
	Sequence 7, Appli	85	85	11:	11:	111	53	28	9	101	10	15	1538	25,	1465	1469	4, 1	127,	1859	1859	6, 2	1551	1551	21,	861	861,	164	86,	1150	115	10,	1246	1246	9

```
APPLICANT: Miller, David J.

APPLICANT: Pease, Larry R.

ITITLE OF INVENTION: Human 1gM Antibodies and Diagnostic and

TITLE OF INVENTION: Therapeutic Uses Thereof Particularly in the Central Nervous

TITLE OF INVENTION: System

FILE REFERENCE: 1199-1-05CIP2

CURRENT FILING DATE: 2001-11-13

PRIOR APPLICATION NUMBER: 09/73,473

PRIOR FILING DATE: 2000-12-05

PRIOR FILING DATE: 2000-05-30

PRIOR APPLICATION NUMBER: 09/32,862

PRIOR APPLICATION NUMBER: 09/32,862

PRIOR APPLICATION NUMBER: 09/32,862

PRIOR APPLICATION NUMBER: 08/79,784

PRIOR APPLICATION NUMBER: 08/79,784

PRIOR PILING DATE: 1999-01-07

PRIOR PILING DATE: 1999-01-07

PRIOR PILING DATE: 1995-08-08

PRIOR FILING DATE: 1994-04-29

NUMBER: 08/236,520

PRIOR FILING DATE: 1994-04-29

NUMBER: 09/236,520

PRIOR PRIOR FILING DATE: 1994-04-29

NUMBER: 09/236,520

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US-10-010-729-9
; Sequence 9, Application US/10010729
; Publication No. US20030185827A1
; GENERAL INFORMATION:
```

Query Match

Similarity

99.3%;

Score 582; DB 14; Pred. No. 1.3e-46;

DB 14;

Length 114;

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TOPOLOGY: linear AA;

MOLECULE TYPE: Light chain - AA
SEQUENCE DESCRIPTION: SEQ ID NO:
US-10-194-801C-4
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US-10-194-801C-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches 112; Conservative
                                                                                                                                                                          Query Match
Best Local (
                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 09/372,425
FILING DATE: AUGUST 11, 1999
ATTORNEY/AGENT INFORMATION:
NAME: Oldenkamp, David J.
REGISTRATION NUMBER: 29,421
REFERENCE/DOCKET NUMBER: 0180.0033
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/194,801C
FILING DATE: 11-Mar-2003
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy Disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Rh Antibody NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Sherie L. Morrison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       61 DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLGQPK 114
61 DRFSGSKSGTSATLGITGLQTGDEADYYCGTWDSSLSAVVFGGGTKLTVLGQPK 114
61 DRFSGSKSGTSATLGITGLQTGDEADYYCGTWDSSLSAVVFGGGTKLTVLGQPK 114
                                                                                                                                                        102;
  79
                  61 DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLGQPK 114
                                                                           19
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                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPRLLIYDITKRPSGIP
DRFSGSKSGTSATLGITGLQTGDEADYYCGTWNSSLSAVLFGGGTEVTVLGQPK 132
                                                                           QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNYVSWYQHFPGTAPKLLIYDNNQRPSGIP
                                                                                               QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: Windows 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Santa Monica
STATE: CA
                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET:
                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (310) 319-5411
TELEFAX: (310) 319-5401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Shapiro & Dupont LLP
STREET: 233 Wilshire Boulevard, Suite 700
                                                                                                                                                                                                                                                                                                                                              LENGTH: 234 amino acids
                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90067
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ramon Montano
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MS Word
                                                                                                                                                                      92.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
                                                                                                                                                        6
                                                                                                                                                                          Score 539; DB 14; Length 234; Pred. No. 2.7e-42;
                                                                                                                                                          Mismatches
                                                                                                                                                        6
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RESULT 3 US-10-779-461-50

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; TYPE: PRT
; ORGANISM: Artificial sequence
; FEATURE:
; OTHER INFORMATION: CM085C11 scFv
US-10-322-673-53
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US-10-322-673-53
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                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: 60/369,877
PRIOR FILING DATE: 2002-04-05
PRIOR APPLICATION NUMBER: 60/384,828
PRIOR FILING DATE: 2002-06-04
PRIOR APPLICATION NUMBER: 60/396,591
PRIOR FILING DATE: 2002-07-18
PRIOR APPLICATION NUMBER: 60/403,370
PRIOR FILING DATE: 2002-08-15
PRIOR FILING DATE: 2002-08-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: MOTTON, Philip A
APPLICANT: MOTTON, ANTIBODIES TO C-MET FOR THE TREATMENT OF CANCERS
TITLE OF INVENTION: ANTIBODIES TO C-MET FOR THE TREATMENT OF CANCERS
FILE REFERENCE: 00980/1
CURRENT FILING DATE: 2004-02-13
CURRENT FILING DATE: 2004-02-13
PRIOR APPLICATION NUMBER: 60/447,073
PRIOR FILING DATE: 2003-02-13
NUMBER OF SEQ ID NOS: 161
SOFTWARE: PATENTIAL 246
TENOMARE: PATENTIAL 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 50, Application US/10779461
Publication No. US20040166544A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS:
SEQ ID NO 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Salcedo et al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 53, Application US/10322673 Publication No. US20030180296A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches 103; Conservative
                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: PF585
CURRENT APPLICATION NUMBER: US/10/322,673
CURRENT FILING DATE: 2002-12-19
PRIOR APPLICATION NUMBER: 60/341,237
PRIOR FILING DATE: 2001-12-20
PRIOR FILING DATE: 2001-12-20
                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: 60/425,737 PRIOR FILING DATE: 2002-11-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: Antibodies that Immunospecifically Bind to TRAIL TITLE OF INVENTION: Receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 24
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: phage display generated human antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: artificial
                                                                                                                                                                                                                                                                                         LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        138 QSVVTQPPSVSAAPGQKVTISCSGSSSNIGNNYVSWYKQLPGTAPKLLIYDNNKRPSGIP 197
  133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLG 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 OSVLTOPPSVSAAPGOKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP 60
                                                                                                                                                                                                                                                                                           243
                         OSVLTOPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP 60
  QSVLTQPPSVSAAPGQKVTISCSGSTSNIGNNYVSWYQQVPGTAPKLLIYDNNKRPSGIP
                                                                                                       91.8%;
91.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92.0%;
92.8%;

 Mismatches

                                                                                       5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 539; DB 16;
Pred. No. 2.9e-42;
                                                                                       Score 538; DB 14;
Pred. No. 3.5e-42;
5; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16;
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RESULT 6
US-10-981-621-53
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CURRENT APPLICATION NUMBER: US/10/981,465

CURRENT FILING DATE: 2004-01-05

PRIOR APPLICATION NUMBER: 60/608,386

PRIOR FILING DATE: 2004-09-10

PRIOR FILING DATE: 2004-09-10

PRIOR APPLICATION NUMBER: 60/468,092

PRIOR APPLICATION NUMBER: 60/495,140

PRIOR FILLING DATE: 2003-08-16

PRIOR APPLICATION NUMBER: 60/495,140

PRIOR APPLICATION NUMBER: 60/396,877

PRIOR FILLING DATE: 2002-04-05

PRIOR PRILING DATE: 2002-04-05

PRIOR PRILING DATE: 2002-04-05

PRIOR APPLICATION NUMBER: 60/369,877

PRIOR FILLING DATE: 2002-04-05

PRIOR APPLICATION NUMBER: 60/394,828

PRIOR APPLICATION NUMBER: 60/394,591

PRIOR APPLICATION NUMBER: 60/396,591

PRIOR APPLICATION NUMBER: 60/396,591

PRIOR PILLING DATE: 2002-07-18

PRIOR PILLING DATE: 2002-07-18

PRIOR PILLING DATE: 2002-07-18

PRIOR PILLING DATE: 2002-07-18

PRIOR APPLICATION NUMBER: 60/403,370

PRIOR PILLING DATE: 2002-07-18

PRIOR APPLICATION NUMBER: 60/425,737

PRIOR PILLING DATE: 2002-08-18

PRIOR PILLING DATE: 2002-11-13

PRIOR PILLING DATE: 2002-08-18

PRIOR PILLING DATE: 2002-11-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 91.8%; Score 538; DB 18; Length 243; Best Local Similarity 91.9%; Pred. No. 3.5e-42;
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                     APPLICANT: Salcedo et al.

TITLE OF INVENTION: Antibodies that Immunospecifically Bind to TRAIL

TITLE OF INVENTION: Receptors

FILE REFERENCE: PF585P1D1

CURRENT APPLICATION NUMBER: US/10/981,621

CURRENT FILING DATE: 2004-11-05

PRIOR APPLICATION NUMBER: 60/608,386

PRIOR FILING DATE: 2004-09-10
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APPLICATION NUMBER: PCT/US04/013900
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; PRIOR FILING DATE: 2004-05-05
; PRIOR APPLICATION NUMBER: 60/468,092
; PRIOR FILING DATE: 2003-05-06
; PRIOR PILING DATE: 2003-05-06
; PRIOR APPLICATION NUMBER: 60/495,140
PRIOR APPLICATION NUMBER: 10/322,673
; PRIOR-FILING DATE: 2002-12-19
; PRIOR APPLICATION NUMBER: 60/369,877
PRIOR FILING DATE: 2002-04-05
; PRIOR APPLICATION NUMBER: 60/364,828
PRIOR APPLICATION NUMBER: 60/396,591
PRIOR PILING DATE: 2002-07-18
; PRIOR PILING DATE: 2002-07-18
; PRIOR PILING DATE: 2002-07-18
; PRIOR PILING DATE: 2002-08-15
                                                    CURRENT APPLICATION NUMBER: US/10/981,673
; CURRENT FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: 60/608,386
; PRIOR APPLICATION NUMBER: 60/608,386
; PRIOR PILING DATE: 2004-09-10
; PRIOR APPLICATION NUMBER: PCT/US04/013900
; PRIOR FILING DATE: 2003-05-06
; PRIOR APPLICATION NUMBER: 60/468,092
; PRIOR APPLICATION NUMBER: 60/469,140
; PRIOR. APPLICATION NUMBER: 10/322,673
; PRIOR APPLICATION NUMBER: 10/322,673
; PRIOR APPLICATION NUMBER: 60/369,877
; PRIOR APPLICATION NUMBER: 60/369,877
; PRIOR APPLICATION NUMBER: 60/398,591
; PRIOR APPLICATION NUMBER: 60/396,591
; PRIOR APPLICATION NUMBER: 60/403,370
; PRIOR FILING DATE: 2002-07-18
; PRIOR APPLICATION NUMBER: 60/403,370
; PRIOR FILING DATE: 2002-01-19
; PRIOR FILING DATE: 2002-07-18
; PRIOR APPLICATION NUMBER: 60/403,370
; PRIOR FILING DATE: 2002-01-19
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; OTHER INFORMATION: CM085C11 8CFV US-10.981-621-53
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US-10-981-673-53
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Publication No. US20050214207A1
GENERAL INFORMATION:
APPLICANT: Salcedo et al.
TITLE OF INVENTION: Antibodies that Immunospecifically
TITLE OF INVENTION: Receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 91.8%; Score 538; DB 18; Best Local Similarity 91.9%; Pred. No. 3.5e-42; Matches 102; Conservative 5; Mismatches 4;
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Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 76
SEQ ID NO 53
LENGTH: 243
TYPE: PRI
ORGANISM: Artificial sequence
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Prior Application SEQ ID NOS: 76
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                                  data removed -
                                      See File Wrapper or PALM
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RESULT 8
US-10-981-691-53
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LENGTH: 243
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LENGTH: 243
TYPE: PRT
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Best Local S
Matches 102
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Antibodies that Immunospecifically Bind TITLE OF INVENTION: Receptors FILE REFERENCE: PF855P1D3 CURRENT APPLICATION NUMBER: US/10/981,691 CURRENT FILING DATE: 2004-11-05 PRIOR APPLICATION NUMBER: 60/608,386 PRIOR PILING DATE: 2004-09-10 PRIOR PILING DATE: 2004-09-05 PRIOR APPLICATION NUMBER: 60/468,092 PRIOR APPLICATION NUMBER: 60/468,092
                                                                                                                                                                                                                                                                                                                                               Remaining Prior Application data removed - See File Wrapper or PALM NUMBER OF SEQ ID NOS: 76
                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: 60/425,737 PRIOR FILING DATE: 2002-11-13
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                                                                                                                                                                                                                                                                      ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                         FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 2002-06-04
APPLICATION NUMBER: 60/396,591
FILING DATE: 2002-07-18
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APPLICATION NUMBER: 60/495,140
FILING DATE: 2003-08-15
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FILING DATE: 2002-04-05
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FILING DATE: 2002-12-19
                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 60/403,370
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Similarity 91.9%;
02; Conservative
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                                                                   QSVLTQPPSVSAAPGQKVTISCSGSTSNIGNNYVSWYQQVPGTAPKLLIYDNNKRPSGIP 192
                                                                                            Q$VLTQPP$V$AAPGQKVTI$C$G$$$NIGNNFV$WYQQLPGTAPXLLIYDITXRP$GIP
DRFSGSKSGTSATLGITGLQTGDEADYYCGTWDSSLSALVFGGGTKVTVLG 243
                   DRFSGSKSGTSATIGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLG 111
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No. US20050214208A1
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91.9%;
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Pred. No. 3.5e-42;
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                                                                                                                                          Score 538; DB 18;
Pred. No. 3.5e-42;
5; Mismatches 4;
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APPLICANT: Kathy Manchulencho
APPLICANT: Raffaella Raggioni
APPLICANT: Glorgio Senaldi
APPLICANT: Glorgio Senaldi
APPLICANT: Glorgio Senaldi
APPLICANT: Olaojuan Jane Su
TITLE OF INVENTION: ANTIBODIES DIRECTED TO TUMOR NECROSIS
TITLE OF INVENTION: FACTOR AND USES THEREOF
FILE REFERENCE: ABGENIX.073A
CURRENT APPLICATION NUMBER: US/10/727,155
CURRENT FILING DATE: 2003-12-02
PRIOR APPLICATION UMBER: 60/430729
PRIOR PILING DATE: 2002-12-02
NUMBER OF SEQ ID NOS: 320
NUMBER OF SEQ ID NOS: 320
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US-10-727-155-309
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US-10-727-155-311
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                                                                                                                                                                                                                                                                                                                                                                   Sequence 311, Applic Publication No. US20 GENERAL INFORMATION:
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Best Local Similarity
            APPLICANT: Palaniswami Rathanaswami
APPLICANT: Craig Pigott
APPLICANT: Meina Liang
APPLICANT: Rozanne Lee
APPLICANT: Kathy Manchulencho
APPLICANT: Kaffaella Faggioni
APPLICANT: Giorgio Senaldi
APPLICANT: Qiaojuan Jane Su
TITLE OF INVENTION: FACTOR AND USES THEREOF
TITLE OF INVENTION: FACTOR AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: John S. Babcook
APPLICANT: Jaspal S. Kang
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                                                                                                                                                                                                                                                                                                                            APPLICANT: John S. APPLICANT: Jaspal
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TYPE: PRT
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REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     104;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVL 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  61 DRFSGSKSGTSATLGITGLQTGDEADYYCGTWDSSLSAGVFGGGTKLTVL 110
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNYVSWYQQLPGTAPKLLIYDNNKRPSGIP
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Xiao Feng
Scott Klakamp
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Rozanne Lee
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                                                                                                                                                                                                                                                                                                                              Jaspal S. Kang
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  ABGENIX.073A
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Pred. No. 1.9e-42;
1; Mismatches 5;
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US-10-293-418-1260
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NUMBER OF SEQ ID NOS: 3239
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 1260
LENGTH: 258
TYPE: PRT
ORGANISM: Homo Bapiens
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US-09-880-748-1260
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; ORGANISM: Homo sapiens
US-10-727-155-311
Sequence 1260, Application US/10293418
Publication No. US20030223996A1
GENERAL INFORMATION:
APPLICANT: Ruben et al.
TITLE OF INVENTION: Antibodies that I
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SOFTWARE: FastSEQ for Windows Version
SEQ ID NO 311
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Best Local :
                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR FILING DATE: 2001-03-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/880,748
CURRENT FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/212,210
PRIOR FILING DATE: 2000-06-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Ruben et al.
TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
FILE REFERENCE: PF523
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 60/240,816 PRIOR FILING DATE: 2000-10-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/10/727,155
CURRENT FILING DATE: 2003-12-02
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                                                                                                                                                                          DRFSGSKSGTSATLGITGLQTGDEADYYCGAWDSSLSAYVFGSGTKLTVLG 258
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    Antibodies that Immunospecifically Bind BLyS
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92.8%;
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Pred. No. 1.9e-42;
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Pred. No. 8.9e-42;
                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: 60/212,210
PRIOR PRIOR PRIOR POLICATION NUMBER: 60/240,816
PRIOR FILING DATE: 2000-10-17
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILING DATE: 2001-03-16
PRIOR PILING DATE: 2001-03-17
PRIOR PILING DATE: 2001-03-21
PRIOR FILING DATE: 2001-03-21
PRIOR FILING DATE: 2001-03-21
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CURRENT APPLICATION NUMBER: US/10/293,418
CURRENT APPLICATION NUMBER: 60/331,469
PRIOR APPLICATION NUMBER: 60/331,469
PRIOR FILING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR FILING DATE: 2001-12-19
PRIOR APPLICATION NUMBER: 09/880,748
PRIOR APPLICATION NUMBER: 09/890,748
PRIOR FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: 60/293,499
PRIOR FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR APPLICATION NUMBER: 60/277,248
PRIOR FILING DATE: 2001-03-21
PRIOR FILING DATE: 2001-03-21
PRIOR FILING DATE: 2001-03-16
PRIOR FILING DATE: 2001-03-16
PRIOR FILING DATE: 2001-03-16
                                                 Query Match
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SEQ ID NO 1246
LENGTH: 248
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SEQ ID NO 1260
LENGTH: 258
TYPE: PRT
    Matches 104;
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/09/880,748
CURRENT FILING DATE: 2001-06-15
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PRIOR APPLICATION NUMBER: 60/212,210
PRIOR FILING DATE: 2000-06-16
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PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/240,816
                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 3239
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Ruben et al.
                                                                                                                         ORGANISM: Homo
                             Local
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                             Similarity
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    Conservative
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                      91.0%;
93.7%;
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Score 533; DB 10;
Pred. No. 1.1e-41;
1; Mismatches 6;
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Pred. No. 8.9e-42;
1; Mismatches 7;
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QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP 60

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; NUMBER OF SEQ ID NOS: 76
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 10
; LENGTH: 112
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-269-805-10
                                                                                                                                                                                        US-10-269-805-10

Sequence 10, Application US/10269805

Publication No. US20030124129A1

GENERAL INFORMATION:

APPLICANT: OLINER, JONATHAN D.

TITLE OF INVENTION: ANGIOPOISTIN-2 SPECIFIC BINDING AGENTS

FILE REFERENCE: A-722

FILE REFERENCE: A-722

CURRENT APPLICATION NUMBER: US/10/269,805

CURRENT FILING DATE: 2002-10-10

PRIOR APPLICATION NUMBER: US 60/328,604

PRIOR PILING DATE: 2001-10-11

PRIOR FILING DATE: 2001-10-11
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CURRENT APPLICATION NUMBER: US/10/293,418
CURRENT FILING DATE: 2002.11-27
PRIOR APPLICATION NUMBER: 60/31,469
PRIOR FILLING DATE: 2001-11-16
PRIOR APPLICATION NUMBER: 60/340,817
PRIOR FILLING DATE: 2001-12-19
PRIOR PILLING DATE: 2001-05-19
PRIOR FILLING DATE: 2001-06-18
PRIOR FILLING DATE: 2001-05-25
PRIOR FILLING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR APPLICATION NUMBER: 60/277,379
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR FILLING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR APPLICATION NUMBER: 60/276,248
PRIOR PILLING DATE: 2001-03-16
PRIOR PILLING DATE: 2001-03-16
PRIOR PILLING DATE: 2001-03-16
PRIOR PILLING DATE: 2000-10-17
PRIOR PILLING DATE: 2000-10-17
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; LENGTH: 248
; TYPE: PRT
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Publication No. US20030223996A1
GENERAL INFORMATION:
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Best Local 9
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TITLE OF INVENTION: Antibodies that Immunospecifically Bind BLyS
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Query Match

90.8%; Score 532; DB 14; Length 112;
Best Local Similarity 92.9%; pred. No. 5.7e-42;
Matches 104; Conservative 2; Mismatches 4; Indels 2; Gaps 1

Qy

1 QSVLTQPPSYSAAPQKVTISCSGSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP 60

1 QSVLTQPPSYSAAPQKVTISCSGSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP 60

1 QSVLTQPPSYSAAPQKVTISCSGSNIGNNFVSWYQQLPGTAPKLLYVDNNKRPSGIP 60

1 QSVLTQPPSYSAAPQKVTISCSGSNSNIGNNFVSWYQQLPGTAPKLLYVDNNKRPSGIP 60

Qy

61 DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSA-VVFGGGTKLTVL 110

Qb

61 DRFSGSKSGTSATLGITGLQTGDEADYYCGTWDSSLSAABVVFGGGTKLTVL 112

Db

61 DRFSGSKSGTSATLGITGLQTGDEADYYCGTWDSSLSAABVVFGGGTKLTVL 112
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Search completed: October 13, 2005, 03:02:48
Job time : 111.575 secs

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Result
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/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
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US-09-240-274-62
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US-09-315-574-3
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Sequence 4, Appli Sequence 18, Appl Sequence 18, Appl Sequence 18, Appl Sequence 2, Appli Sequence 2, Appli Sequence 62, Appli Sequence 61, Appli Sequence 61, Appli Sequence 5, Appli Sequence 5, Appli Sequence 4, Appli Sequence 4, Appli Sequence 7, Appli Sequence 42, Appli Sequence 43, Appli Sequence 44, Appli Sequence 45, Appli Sequence 45, Appli Sequence 44, Appli Sequence 45, Appli Sequence 
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	Match ocal	REFERENCE/DOC TELECOMMUNICATI TELEPAN: (31 TELEPAX: (31 FFORMATION FOR S SEQUENCE CHARAC LENGTH: 234 TYPE: amino STRANDENMESS: TOPOLOGY: 11 MOLECULB TYPE:	STATES CA STATES CA COUNTRY: USA ZIP: 90067 ZIP: 90067 ZIP: 90067 ZIP: PORE MEDIUM TYPE: PLOPP COMPUTER READABLE FORE MEDIUM TYPE: PLOPP COMPUTER: IEM PC CO OPERATING SYSTEM: 1 SOFTWARE: MS WOR'D CURRENT APPLICATION NUMBER: PILING DATE: AUGUST CLASSIFICATION UNTER CLASSIFICATION UNTER CLASSIFICATION UNTER CLASSIFICATION UNTER CLASSIFICATION NUMBER: PILING DATE: 11FORM ATTORNEY/AGENT INFORM NAME: Oldenakmp, DA REGISTRATION NUMBER	- W & W & _ LE LE	5 5 5
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	ati	IT NUMB I INFOR 0) 788-5 788-5 ID NO RISTIC LING AC SIG Single sar Light C	FORM: FORM: Coppy Ccom Ccom N DATA: ER: Gust 135 DATA: ER: ER: DATA: DATA: ER:	<b>⊑00</b> 3	
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	40 44	REFERENCE/DOCKET NUMBER: 51 TELECOMUNICATION INFORMATION: TELEPHONE: (310) 788-5000 TELEPHONE: (310) 788-5100 TELEPAX: (310) 788-5100 TELEPAX: (310) 788-5100 TOMATION FOR SEQ ID NO: 4: SEQUENCE CHARACTERISTICS: LENGTH: 234 amino acids TYPE: amino acid TYPE: amino acid STRANDEDNESS: single TOPOLOGY: linear MOLECULE TYPE: Light chain372-425A-4	STATES CA COUNTRY: LOS ANGELES STATES CA COUNTRY: USA ZIP: 90067 COMPUTER READABLE FORM: MEDIUM TYPE: Ploppy Disk COMPUTER: IBM PC compatible OPERATING SYSTEM: Windows 98 SOFTWARE: MS WOR'd CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/372,4 PRICH APPLICATION DATA: PRICH APPLICATION DATA: PRICH APPLICATION DATA: PRICH APPLICATION UNUMBER: PRICH APPLICATION DATA: PRICH APPLICATION DATA: PRICH APPLICATION DATA: PRICH ODATE: APPLICATION NUMBER: 29,421 REGISTRATION NUMBER: 29,421	ar Koll 6	0 4 0 4 u 4 4 u 1 4 u 1 0 4 0 1 u u
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; MOLECULE TYPE: protein
US-09-025-769B-18
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Matches
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                Sequence 18, Application US/09490070A Patent No. 6696248 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (212)596-909
INFORMATION FOR SEQ ID NO:
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Ge, LAWRING MORONEY, SIMON
APPLICANT: Moroney, Simon
APPLICANT: Plueckthun, Andreas
APPLICANT: Plueckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
TITLE OF STOHENCES: 373
                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 112 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: James R. Haley, Jr., Esq. REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: MORPHO/5
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/025,769B
FILING DATE: 18-FEB-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: EP 95 11 3021.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC COmpatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 18-AUG-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                       y Match 88.4%;
Local Similarity 87.5%;
hes 98; Conservative !
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
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APPLICANT: Knappik, Achim
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                                                                                                                                                         DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLGQ 112
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                                                                                                                                       DRFSGSKSGTSASLAITGLQSEDEADYYCATWDDSLSGVVFGGGTKLTVLGQ 112
                                                                                                                                                                                                       QSVLTQPPSVSGAPGQRVTISCSGSSSNIGNNYVSWYQQLPGTAPKLLIYDNNKRPSGVP
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1251 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                  112 amino acids
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Pack, Peter
Ilag, Vic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (212)596-9000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EP 95 11 3021.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18:
                                                                                                                                                                                                                                                                       Score 518; DB 3;
Pred. No. 6.4e-44;
5; Mismatches 9
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                                                                                                                                                                                                                                                                                                           Length 112;
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Matches
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TOPOLOGY: Ilnear;
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO:
US-09-490-070A-18
                                                                                                                                                                                                                                                             Sequence 18, Application Patent No. 6706484
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: COLIN G. Sandercock, Bsq.
REGISTRATION UNMBER: 31,298
REFERENCE/DOCKET NUMBER: 37629-0005
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 912-2000
TELEPAX: (202) 912-2020
INFORMATION FOR SEQ ID NO: 18:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Versi
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/490,070A
FILING DATE: 24-Jan-2000
PRIOR APPLICATION NUMBER: EP 95 11 3021.0
FILING DATE: 18-AUG-1995
TILING DATE: 18-AUG-1995
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MEDIUM TYPE: Floppy disk
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Plueckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
NUMBER OF SEQUENCES: 373
Moroney, Simon
PlueCkthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
NUMBER OF SEQUENCES: 373
CORRESPONDENCE ADDRESS:
ADDRESSEE: James F. Haley, Jr., Esq. c/o Fish & Neave
STREET; 1251 Avenue of the Americas
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                        APPLICANT: Knappik, Achim
                                                                                                                                                                                                                                                                                                                                                                                                                                             61 DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLGQ 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              QSVLTQPPSVSAAPGQKYTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP 60
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TYPE: amino acid
STRANDEDNESS: <Unknown>
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                                                                                                                                                                                                                                                                                                                                                                                                                    DRFSGSKSGTSASLAITGLQSEDEADYYCATWDDSLSGVVFGGGTKLTVLGQ 112
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White & McAuliffe
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Ilag, Vic
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Pred. No. 6.4e-44;
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US-09-490-324-18
Sequence 18, Application US/09490324; Patent No. 6828422; GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Pateentin Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: protein SEQUENCE DESCRIPTION: S
                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                          Moroney, Simon
Plueckthun, Andreas
TITLE OF INVENTION: Protein/(Poly)peptide libraries
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FILING DATE: 24-Jan-2000
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/025,769B
FILING DATE: 18-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.30 (RPO) CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 QSVLTQPPSVSAAPGQXVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                              ADDRESSEE: James F. Haley, Jr., Esq. c/o Fish & Neave STREET: 1251 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QSVLTQPPSVSGAPGQRVTISCSGSSSNIGNNYVSWYQQLPGTAPKLLIYDNNKRPSGVP
APPLICATION NUMBER: US/09/490,324
                                                                                                                                                                             STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DRFSGSKSGTSASLAITGLQSEDEADYYCATWDDSLSGVVFGGGTKLTVLGQ 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: <Unknown>
                                                                                                                                         ZIP: 10021
                                                                                                                                                           COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: James F. Haley, Jr., Esq. REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: MORPHO/5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 112 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: James F.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (212)596-9000
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                                                                                                                                                                                                                                                                                                                                                                          Pack, Pete
                                                                                                                                                                                                                                                                                                                                                                                                                   Knappik, Achim
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oney, Simon
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Pred. No. 6.4e-44;
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Best Local &
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                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: PSTENT: Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/199,911
FILING DATE: 22-PEB-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/065,517
FILING DATE: 21-MAY-1993
PRIOR APPLICATION DATA:
                                                                        PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Kobrin, Barry J.
APPLICANT: Haspel, Martin V.
TITLE OF INVENTION: Tumor Associated Monoclonal Antibody
NIMBER OF INVENTION: 123AV16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                            · COUNTRY:
                                                                                               APPLICATION NUMBER: US 07/636,179 FILING DATE: 31-DEC-1994
  APPLICATION NUMBER:
                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Rockville STATE: Maryland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: MORPHO/5 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 24-Jan-2000
PRIOR APPLICATION DATA:
APPLICATION UMMBER: US/09/025,769
FILING DATE: 18-FEB-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                              Maryland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/08199911
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: <Unknown>
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TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: EP 95 11 3021.0 FILING DATE: 18-AUG-1995
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TELEFAX: (212)596-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: James F. Haley, Jr., REGISTRATION NUMBER: 27,794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              POPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1330 Piccard Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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                                      25-JAN-1989
                                                                                                                                                                                                                                                                                                                                                    Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        88.4%;
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                                                      US 07/302,155
US 06/697,078
                                                                                                                                                                                                                                                  US/08/199,911
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Pred. No. 6.4e-44;
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                                                                                                                                                                                                                                                                                           Version #1.25
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FILING DATE: 31-JAN-1985
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0.
FILING DATE: 31-JAN-1984
ATTORNEY/AGENT INFORMATION:

US 06/575,533

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RESULT 7
US-09-240-274-62
RESULT 8
US-08-983-607-31
; Sequence 31, Application US/08983607
; Patent No. 6140470
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: Siegel, Donald L.
APPLICANT: Siegel, Donald L.
TITLE OF INVENTION: Rh(D)-BINDING PROTEINS AND MAGNETICALLY ACTIVATED CELL
TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF
FILE REFERENCE: 09596-42U2
CURRENT FILING NATE: 1999-01-29
EARLIER APPLICATION NUMBER: 60/081,380
EARLIER APPLICATION NUMBER: 60/081,380
EARLIER FILING DATE: 1998-04-10
EARLIER FILING DATE: 1998-04-10
EARLIER FILING DATE: 1996-10-11
NUMBER OF SEQ ID NOS: 224
SOFTWARE: Patentin Ver: 2.0
                                                                                                                                                                                                                                                                                                        Matches 101; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 62, Application US/09240274
Patent No. 6255455
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 301-977-0847
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
OTHER INFORMATION: anti-Rh(D) chain NO2
                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 114
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 110 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: 301-258-5200
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE:
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Local Similarity 89.1%;
                                                                                                                                                        64 SGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSA----VVFGGGTKLTVLG 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          61 DRFSGSKSGTSATLGITGLQTGDEADYYCGTWDTRLRAGVFGGGTKLTVL 110
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                                                                                                                                                                                                                                             4 LTQPPSVSAAPGQXVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIPDRF 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVL 110
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                                                                                                                                  SGSKSGTSATLGITGLQTGDEADYYCGTWDSSLSAGRVRRMFGGGTKLTVLG 114
                                                                                                                                                                                                                       LTQPPSVSAAPGQKVTISCSGSSSNIGNNYVSWYQQLPGTAPKLLIYDNNKRPSGIPDRF
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90.2%;
                                                                                                                                                                                                                                                                                                     Score 515; DB 3; Length 114;
Pred. No. 1.3e-43;
2; Mismatches 5; Indels
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US-08-345-321-4
                                       RESULT 9
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APPLICANT: Alan Garen
APPLICANT: Xiaohong Cai
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OPERATING SYSTEM: MS DOS
SOFTWARE: WOR'D PROCESSING
SOFTWARE: WOR'D PROCESSING
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/983
FILING DATE: April 27, 1998
CLASSIFICATION DATA:
APPLICATION NUMBER: PCT/IB96/
PILING DATE: June 28, 1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MARY M. KLINSKY
N
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
NAME/KEY: light chain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
DESCRIPTION: polypeptide
ORIGINAL SOURCE:
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MEDIUM TYPE: 3.5" 1.44 Mb diskette
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ORGANISM: nized with autologous tumor cells)
INDIVIDUAL 1801ATE: peripheral blood lymphocytes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: 203-773-9544
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Y MALCH
Local Similarity 90.2%;
les 101; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LIBRARY: DM414 scFv antibodies obtained from LIBRARY: fUSE5 fusion phage construct CLONE: V73
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STATE: Connecti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: sir
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER:
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                                                                                                                                                      61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 513.5; DB 3; Length 112;
Pred. No. 1.8e-43;
3; Mismatches 7; Indels 1
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                                                                                                                                                              112
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; Sequence 4, Application US/08345321; Patent No. 5914109; GENERAL INFORMATION:

APPLICANT:

ZOLLA-PAZNER, Susan

APPLICANT: GORNY, MITOSIAV K.
TITLE OF INVENTION: METERCHYBRIDOMAS PRODUCING HUMAN
TITLE OF INVENTION: MONOCLONAL ANTIBODIES TO HIV-1

NUMBER OF SEQUENCES:

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US-09-240-274-61
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                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No. 6255455
GENERAL INFORMATION:
                                                   SEQ ID NO 61
LENGTH: 109
                                                                                                                     APPLICANT: Siegel, Donald L.

TITLE OF INVENTION: Rh (D)-BIDING PROTEINS AND MAGNETICALLY ACTIVATED CELL

TITLE OF INVENTION: SORTING METHOD FOR PRODUCTION THEREOF

FILE REFERENCE: 09596-42U2

CURRENT FILING DATE: 1999-01-29

EARLIER APPLICATION NUMBER: 60/081,380

EARLIER APPLICATION NUMBER: 60/080

EARLIER FILING DATE: 1996-10-11

EARLIER FILING DATE: 1996-10-11

NUMBER: 05-20-1

NUMBER: 05-20-1
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TELEX: 248633
INFORMATION FOR SEQ ID NO:
                                                                                                       SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Browdy, Roger L.
REGISTRATION NUMBER: 25,618
REFERENCE/DOCKET NUMBER: ZOLI
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
TELEPAX: 202-737-3528
ORGANISM: Homo sapiens
                                TYPE: PRT
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LENGTH: 132 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/345,321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: protein
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Browdy and Neimark
STREET: 419 Seventh Street, N.W., Suite 300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       h 84.4%; Score 494.5; DB 2
Similarity 88.3%; Pred. No. 1.6e-41;
98; Conservative 2; Mismatches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSA-VVFGGGTKLTVL 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               QSVLTQPPSVSAAPGQKVT1SCSGSSSN1PNNYVLWYQQFPGTAPKLL1YGNNKRPSG1P
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Query Match 84.1%;
Best Local Similarity 87.9%;
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; OTHER INFORMATION: anti-Rh(D) chain N01
US-09-240-274-61
                                                                                                                                                          US-08-665-202-36
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                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION: 424

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/000,238

PILING DATE: 14-JUN-1995

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/000,250

PILING DATE: 15-JUN-1995

ATTORNEY/AGENT INFORMATION:
NAME: HUNCEY, Tom
REGISTRATION NUMBER: 38,498
REFERENCE/DOCKST NUMBER: 02307E-061410

TELECOMMUNICATION INFORMATION:
TELECHONE: (415) 576-0200
                                                                               Matches
                                                                                                                                                                             MOLECULE TYPE:
                                                                                                  Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent No. 5977322
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                      TELEPAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
LENGTH: 111 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Marks, James D.
APPLICANT: Schier, Robert
TITLE OF INVENTION: Two S977322el High Affinity Human Antibodies to
TITLE OF INVENTION: Two Antigens
NUMBER OF SEQUENCES: 141
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/665,202
APPLICATION NUMBER: US/08/665,202
FILING DATE: 13-UN-1996
                                                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       3 LTQPPSVSAAPGQKVTISCSGSSSNIDSNYVSWYQQLPGTAPKLL1FDNYRRPSGIPDRP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              4 LTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIPDRF 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  94; Conservative
                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94111-3834
                                                                                                                                                                                                                                       amino acid
QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SGSKSGTSATLGITGLOTGDEADYYCXTWDSSLSAVVFGGGTKLTVL 110
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                                                                                                                                                                                                  linear
                                                                                               82.9%;
                                                                                               Score 486; DB 2;
Pred. No. 9.1e-41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 493; DB 3;
Pred. No. 1.8e-41;
                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                  DB 2; Length 111;
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                                                                             Indels
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DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLG 111

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RESULT 12
US-09-315-574-36
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RESULT 13
US-08-665-202-5
; Sequence 5, Application US/08665202
; Patent No. 5977322
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                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (415) 576-030
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Hunter, Tom
REGISTRATION NUMBER: 38,498
REGISTRATION NUMBER: 02307E-061411
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 60/000,250
FILING DATE: 15-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/665,202
FILING DATE: 13-JUN-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/000,238
FILING DATE: 14-JUN-1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Schier, Robert
TITLE OF INVENTION: No. 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US, FILING DATE: 20-MAY-99 CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET:
                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 111 amino acids
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                                                                                                                                    61 DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLG 111
                                                                                                                                                                                                                                                             92; Conservative
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                                                                                                                                                                                    QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNYVSWYQQLPGTAPXLLIYDITKRPSGIP 60
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                                                                                                               DRFSGSKSGTSASLAISGFRSEDEADYYCAAWDDSLSGWVFGGGTKLTVLG 111
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                                                                                                                                                                                                                                                                                                                                                                          linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   No. 6512097el High Affinity Human Antibodies to
                                                                                                                                                                                                                                                                              82.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US/09/315,574
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  36:
                                                                                                                                                                                                                                                         Score 486; DB 4; Length 111;
Pred. No. 9.1e-41;
7; Mismatches 12; Indels
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RESULT 14
US-09-315-574-5
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                                                                                                                                                                                                                                                              Sequence 5, Application US/09315574
Patent No. 6512097
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity Matches 92; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/000,238
FILING DATE: 14-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION UNMBER: US 60/000,250
PILING DATE: 15-JUN-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                               GENERAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Marks, James D.
APPLICANT: Schier, Robert
TITLE OF INVENTION: No. 5977322el High Affinity Human Antibodies to
TITLE OF INVENTION: Tumor Antigens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Hunter, Tom
REGISTRATION NUMBER: 38,498
REFERENCE/DOCKST NUMBER: 02
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/665,202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                     APPLICANT: Marks, James D. APPLICANT: Schier, Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                             NUMBER OF SEQUENCES: 141
CORRESPONDENCE ADDRESS:
ADDRESSES: Majestic, Parsons, Siebert & Hsue P.C.
STREET: Four Embarcadero Center, Suite 1100
                                                                                                                                                             APPLICANT: Schier, Robert
TITLE OF INVENTION: No. 6512097el High Affinity Human Antibodies to
TITLE OF INVENTION: Tumor Antigens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 13-JUN-1996
CLASSIFICATION: 424
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STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY:
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                                     CITY: San Francisco
STATE: California
                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   145
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                                                                                                                                                                                                                                                                                                                                                                                                                       61 DRPSGSKSGTSATIGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLG 111
                                                                                                                                                                                                                                             INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP
94111-4106
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                       USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         <u>ب</u>
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Pred. No. 2.4e-40;
7; Mismatches 12
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RESULT 15
US-09-069-821-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Hunter, Tom
REGISTRATION NUMBER: 38,498
REFERENCE/DOCKET NUMBER: 02307E-061411
TELECOMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPHONE: (415) 576-0300
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 258 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 4
Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 82.9%; Score 486; DB 4; Length 258; Best Local Similarity 82.9%; Pred. No. 2.4e-40; Matches 92; Conservative 7; Mismatches 12; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                 APPLICANT: WHITLOW, MAKE
APPLICANT: LEE, LIHSYNG S.
TITLE OF INVENTION: SINGLE-CHAIN ANTIGEN-BINDING PROTEINS
TITLE OF INVENTION: CAPABLE OF GLYCOSYLATION, PRODUCTION AND USES THEREOF
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
ADDRESSEB: STERNE, KESSLER, GOLDSTEIN & FOX P.L.L.C.
STREET: 1100 NEW YORK AVE., NW, SUITE 600
                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
APPLICANT:
APPLICANT:
                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 60/000,250
FILING DATE: 15-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/665,202
FILING DATE: 13-JUN-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/000,238
FILING DATE: 14-JUN-1995
PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                              COUNTRY:
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SHORR, ROBERT
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Search completed: October 13, 2005, 02:45:19 Job time: 18.5923 secs
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Best Local Similarity
Matches 92; Conserv
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TELEFAX: (202)371-2540
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: KIM, JUDITH U.
REGISTRATION NUMBER: 40,679
REFERENCE/DOCKET NUMBER: 09
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 262 amino acids
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FILING DATE: 23-JUN-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/044,449
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Listing first 45 summaries
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Perfect score:
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GO; GO:0005576; C:extracellular; NAS.
GO; GO:0003823; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR007110; Ig-like.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG LIKE; 1.
Immunoglobulin V region; Signal.
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                                                                                                                               01-JUN-2002 (TrEMBLrel. 21
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Immunglobulin light chain
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01-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleic Acids Res. 12:8407-8414(1984).
                                                                                   Homo sapiens (Human)
Eukaryota; Metazoa;
                                                                                                                                                                                                                                              QSTE63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=85062823; PubMed=6095199;
SEQUENCE
                                                                     Mammalia;
                                                                                                                                                                                                                                                                                            w
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                OSVLTQPPSVSAAPGQKYTISCSGSSSNIGNDYVSWYQQVPGTAPKLLIYDNNKRPSGIP
  FROM N.A.
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                                                                Eutheria;
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                                                                   Primates;
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                                                                                       Chordata;
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25,
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    Last sequence update)
    Last annotation update)
    variable region (Fragment).

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J segment.
By similarity.
                                                                                                                                                                                                      Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 523; DB 1
Pred. No. 2e-43;
                                                                   Catarrhini;
                                                                                       Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                  PRT;
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                                                                   Hominidae;
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RESULT 4
LV1C_HUMAN
ID LV1C_HUMAN
AC P01701;
DT 21-JUL-1986
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                                                                                          SQ TTTT
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Best Local S
Matches 96
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
05-JUL-2004 (Rel. 44, Last annotation updat
15 lambda chain V-I region NEW.
                                                                                                             DOMAIN
MOD RES
DISULFID
NON_TER
                                                                                                                                                                                            Pfam; PF00047; ig; 1.

SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG LIKE; 1.

Bence-Jones protein; Direct protein sequencing;

Immunoglobulin V region; Pyrrolidone carboxylic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; L33985; AAL68704.1; -. HSSP; P01703; 7FAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            type). Subgroups i
the lambda-type.";
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Hall B.L., Murray J.H., Haspel M.
"Establishment, molecular rescue,
                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                             GO; GO:0005576; C:extracellular; NAS. GO; GO:0003823; F:antigen binding; NAS GO; GO:0006955; P:immune response; NAS
                                                                                                                                                                                                                                                                                                                                                                                                                               PIR; A01964; L1HUNW.
HSSP; P01703; 7FAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hoppe-Seyler's Z. Physiol. Chem.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "The complete amino acid sequence of Bence Jones protein type). Subgroups in the variable part of immunoglobulin I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                               InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         - I- MISCELLANEOUS: This is a Bence-Jones protein.
I- SIMILARITY: Contains 1 immunoglobulin-like do
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              anger B., Steinmetz-Kayne M., Hilschmann N.;
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89.1%;
                    86.7%;
                                                                                      MW;
Pred. No. 5e-4
7; Mismatches
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Pred. No. 1e-42;
3; Mismatches
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                                                                                                                                  Ig-like.
Pyrrolidone carboxylic
By similarity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immunoglobulin light chain variable
                                                                                          AAECBCA3C49F2AD3
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                                            Length 111;
                                                                                             CRC64;
                                                                                                                                                                                                          acid.
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                                                                                                                                                              acid
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Similarity

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RESULT 5
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RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan R.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carrinci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carrinci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren B.J., Lu X., Gibbs R.A.,
RA Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Hitting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Karywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Alnes S.J., Marra M.A.
                                                                                               Pfam; PR0765-7, G; 2.

SMART; SM00409; IG; 2.

SMART; SM00407; IGc1; 1.

SMART; SM00406; IGv; 1.

R PROSITE; PS500390; IG MHC; UNKNOWN 1.

PROSITE; PS90290; IG MHC; UNKNOWN 1.
                                                     Query Match
Best Local S
Matches 97
                                                                                                                                                                                                                   InterPro; IPR003599; Ig.
InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig_cl.
InterPro; IPR003597; Ig_MCC.
InterPro; IPR003596; Ig_v.
Pfam; PP07654; Cl-set; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-JUL-2004
05-JUL-2004
05-JUL-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Q6IN99;
                                                                                                                                                                                                                                                                                                                                        Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases. EMBL, BC072392; AAH72392.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                               "Generation and initial analysis of more than 15,000 and mouse cDNA sequences."; Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IGLC2 protein.
Name=IGLC2;
                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                              HSSP; P01842; 1AQK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Jones S.J., Marra M.A.;
                                                                                                                                                                                                                                                                                                                                                                                               TISSUE=Pancreas;
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                                                       97;
                                                                     Similarity
OSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP
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                                                       Conservative
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Primates;
                                                                       85
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27,
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Last annotation update)
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                                                     6,
                                                   Score 507; DB Pred. No. 1.5e 6; Mismatches
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                                                                                                                   90C95D5E87A6BCC1 CRC64;
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                                                                   DB 2;
.5e-41;
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                                                   11,
                                                                                   Length 235;
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Best Local S
Matches 91
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InterPro; IPR003596; Ig-v.
SMART; SM00406; IGv; I-
PROSITE; PS50835; IG_LIXE; 1.
NON TER 1 1 101
SEQUENCE 101 AA; 10374 MW;
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899; Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D., Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K., Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F., Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L., Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E., Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C., Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C., Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C., Brownstein M.J., WcKernan K.J., Malek J.A., Gunaratne P.H., Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W., Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
                                                                                                                                                                                                                                                                                                                    QGCMX4;
QGCMX4;
05-JUL-2004 (TrEMBLrel. 27,
05-JUL-2004 (TrEMBLrel. 27,
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QSIZDS;
01-MAR-2003
                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                             Hypothetical protein.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                               Q6GMX4
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Submitted (AUG-2002) to
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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01-MAR-2003 (TrEMBLrel. 23, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Anti-thyroglobulin light chain variable region (F
                                                                                                                                                                                             TISSUE-Spleen;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SAAPGQKVTISCSGSSSNIGKNYVSWYQQVPGTAPQLLIHDTDKRPSGIPDRFSGSQSGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                QSVLTQPPSVSAAPGQRVTISCSGRSSNIGNSYVSWYQQFPGAAPQLLIYDNDKRPSGIP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              101 AA; 10374 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                 PRELIMINARY;
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90.1%;
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Last annotation update)
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Pred. No. 2e-39;
5; Mismatches
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Best Local S
Matches 92
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PO6888
O1-JAN-1988 (Rel. 06, Created
O1-JAN-1988 (Rel. 06, Last se
05-JUL-2004 (Rel. 44, Last ar
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InterPro; IPR00359; Ig_cl.
InterPro; IPR003597; Ig_cl.
InterPro; IPR003597; Ig_cl.
InterPro; IPR003596; Ig_w.
Pfam; PF00647; Cl-set; I.
Pfam; PF00047; Ig; 2.
SMART; SM00409; IG; 2.
SMART; SM00407; IGcl; 1.
                                                                                                                                                                                                                                                                      SEQUENCE.

MEDLINE=86000126; PubMed=3929803;

MEDLINE=86000126; PubMed=3929803;

Toft K.G., Sletten K., Husby G.;

"The amino-acid sequence of the variable containing amyloid fibril protein EPS (in
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SEQUENCE 23
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                PIR; A24656; L1HUEP.
HSSP; P01703; 7FAB.
GO; GO:0005576; C:extracellular; NAS.
GO; GO:0003823; F:antigen binding; NI
GO; GO:0006955; P:immune response; NI
InterPro; IPR007110; Ig-like.
                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jones S.J., Marra M.A.
                                                                                                                                                     NCBI_TaxID=9606;
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236 AA; 2
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Primates;
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                                                                                                                                                            1 immunoglobulin-like
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence update) annotation updat EPS.
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Pred. No. 3.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               109
                                                                                                                                                                                                                                                                           le region of a carbohydrate-
(immunoglobulin light chain,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               B
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RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA HOpkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Stapleton M.J., Usdin T.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P. Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Mozwan P.J., McKernan K.J., Malek J.A., Gunarathe P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Fahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Krzywinski M.I., Skalska U., Smailus D.E., Schmerch A., Schein J.E.,
RA Jones S.J., Marra M.A.:
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PROSITE; P850835; IG_LIKE; 1.

Amyloid; Direct protein sequencing; Glycoprotein;

Immunoglobulin V recent Ta-11ke
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05-JUL-2004 (TrEMBLrel.
05-JUL-2004 (TrEMBLrel.
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SMART; SM00406; IGv;
                                                 InterPro;
InterPro;
                                                                                                                                                                                           Strausberg R.;
Submitted (JUN-2004) to the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE=Primary B-Cells;
MEDLINE=2238857; PubMed=12477932; DOI=10.1073/pnas.242603899;
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Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hypothetical protein. Homo sapiens (Human).
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                                                                                       EMBL; BC073784; AAH73784.1;
InterPro; IPR003599; Ig.
InterPro; IPR007110; Ig-like
InterPro; IPR003597; Ig c1.
                                                                                                                                                                                                                                     TISSUE=Primary B-Cells;
                                                                                                                                                                                                                                                                 SEQUENCE FROM
                                                                                                                                                                                                                                                                                                                                                                                    Jones S.J., Marra M.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                "Generation and initial analysis
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                                                                                                                                                                                                                                                                                                                                    mouse cDNA sequences.
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90; Conserv
            DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLG
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                                                                                                                                                                                             EMBL/GenBank/DDBJ databases
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Pred. No. 4.1e-38;
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                                                                                                                                                                                                                                                                                                                   99:16899-16903 (2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 556A313E24D5AC73 CRC64;
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RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

RX MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

RA Klausmer R.D., Collins P.S., Wagner L., Shenmen C.M., Schuler G.D.,

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RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mıllahy S.J.,

RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hullyk S.W.,

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RA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

RA Motring R.W., Touchman J.W., Green B.D., Dickson M.C.,

RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

RA Krzyvinski M.I., Skalska U., Smallus D.E., Schmerch A., Schein J.E.,

RA Tones S.T. Marra M.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         밁
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Best Local S
Matches 87
                                                                                  InterPro; IPR007110; Ig-like.
InterPro; IPR003597; Ig_cl.
InterPro; IPR003006; Ig_WHC.
InterPro; IPR003596; Ig_v.
Pfam; PP07654; Cl-set; 1.
SMART; SM00406; IGv; 1.
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SMART; SM00407; IGc1; 1.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 2.
PROSITE; PS00290; IG_MHC; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-OCT-2002 (TrEMBLrel.
01-OCT-2002 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
                                                                                                                                                                                                                                                             EMBL; BC030984; AAH30984.1;
HSSP; P01703; 7FAB.
                                                                                                                                                                                                                                                                                                        Strausberg R.;
Submitted (JUN-2002) to the EMBL/GenBank/DDBJ databases.
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Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hypothetical protein. Homo sapiens (Human).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Generation and initial analysis of more than 15,000 and mouse cDNA sequences."; Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jones S.J., Marra M.A.;
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                     M00406; IGv; 1.
PS50835; IG_LIKE; 2.
PS00290; IG_MHC; UNKNOWN_1.
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                                                                                                                                                                                                                                                                                                                                                                                                            FROM N.A.
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76.3%;
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Last annotation updat
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Pred. No. 3e-37;
1; Mismatches
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RESULT 11
LV1B HUMAN
ID LV1B H
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DT OF
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                                                                                                                                                                                                                                                                       GO; GO:0005576; C:extracellular; NAS.
GO; GO:0003523; F:antigen binding; NAS.
GO:0003525; P:immune response; NAS.
InterPro; IPR007110; Ig-like.
InterPro; IPR007110; Ig-like.
InterPro; IPR0047; ig, 1.
SMART; SM00406; IGv; 1.
SMART; SM00406; IGv; 1.
Bence-Jones protein; Direct protein seque Immunoglobulin V region; Pyrrolidone carb DOMAIN

1 106 Ig-like.
Q6GMX3;
05-JUL-2004
05-JUL-2004
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21-JUL-1986 (Rel. (
05-JUL-2004 (Rel. 4
1g lambda chain V-1
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SEQUENCE
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P01700;
21-JUL-1986
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                                                                                                                                                                                                                                                                                                                                                                                                                         J. Biol. Chem. 245:4475-4487(1970).
-!- MISCELLANEOUS: This is a Bence-Jones protein.
-!- SIMILARITY: Contains 1 immunoglobulin-like domain
                                 Q6GMX3
                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE-71103824; PubMed=5532227; Shinoda T., Titani K., Putnam F.W.; Shinoda T., Titani K., Putnam F.W.; "Amino acid sequence of human lambda chains. II. Chymotryptic peptides and sequence of protein Ha.";
                                                                                                                                                                                                                                                                                                                                                                                                     PIR; A01963; L1HUHA.
HSSP; P01703; 7FAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                     PDRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVL 110
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                                                                                        PDRFSGSKSGTSASLAISGLRSEDEAHYHCAAWDYRLSAVVFGGGTQLTVL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DRFSGSKSGTSASLAISGLRSEDEADYYCAAWDDSLSVHVVFGGGTKLTVLSQPK 134
                                                                                                                                                                                                                              112 AA; 11896 MW;
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                                                                                                                                                                                 Conservative
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                                  PRELIMINARY;
                                                                                                                                                                                                                                                                     V region; Pyrrolidone carboxylic 1 106 Iq-like
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76.6%;
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76.5%;
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Pred. No. 5.1e-35;
9; Mismatches 16;
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Pred. No. 4.:
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                                                                                                                                                                                                                             8D73378F3F5CD039 CRC64;
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(TrEMBLrel. (TrEMBLrel.

27, 27,

Last sequence update)

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X KIRLUINE-22388527; PubMed=12477932; DOI=10.1073/pnas.242603899;

X Klausher R.D., Collins F.S., Wagner L.H., Derge J.G.,

X Klausher R.D., Collins F.S., Wagner L. Shemmen C.M., Schuler G.D.,

X Altschul S.F., Zeeberg B., Buerow K.H., Schaefer C.F., Bhat N.K.,

X Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

X Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

X Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

X Hopkins R.F., Jordan K., Farmer A.A., Rubin G.M., Hong L.,

X Hopkins R.F., Jordan K., Farmer A.A., Rubin G.M., Hong L.,

X Hopkins R.F., Jordan K., Farmer A.A., Rubin G.M., Hong L.,

X Hopkins R.F., Jordan K., Farmer A.A., Rubin G.M., Hong L.,

X Hopkins R.F., Jordan R.B., Toshiyuki S., Carninci P., Prange C.,

X Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

X Raha S.S., Loquellano N.A., Peters G.J., Malek J.A., Gunarathe P.H.,

X Raha S.S., Loquellano N.A., Peters G.J., Malek J.A., Gunarathe P.H.,

X Richards S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.M.,

X Hilalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

X Pahey J., Helton B., Young A.C., Shevchenko Y., Bouffard G.G.,

X Pahey J., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

X Holden R.F., Jordan B., Schmutz J., Myers R.M., Butterfield Y.S.,

X Topes G. J. Mayra M. A.,

X To
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Best Local S
Matches 86
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A Strausberg R.;
Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases.

EMBL, BEO73770; AAH73770.1; -.

R InterPro; IPR003599; IG.
R InterPro; IPR003597; IG-11.
R InterPro; IPR003597; IG-11.
R InterPro; IPR003596; IG-MHC.
R Pfam; PP07654; C1-set; 1.
R Pfam; PP0047; IG; 2.
R SMART; SM00409; IG; 2.
R SMART; SM00409; IG-11.
R SMART; SM00407; IG-11.
R SMART; SM00409; IG-11.
R SMART; SM00409; IG-11.
R PROSITE; PS50835; IG-LIKE; 2.
Q96E61
Q96E61;
01-DEC-2001 (
01-DEC-2001 (
01-MAR-2004 (
Hypothetical
Homo sapiens
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SEQUENCE 2
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TISSUE=Spleen;
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Mammalia; Eutheria; Primates;
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Homo sapiens (Human).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    QSVLTQPPSVSGAPGQGVTISCTGSSSNIGAGYGVHWYQQLPGTAPKLLISNNSNRPSGV
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236 AA; 2
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   protein. (Human).
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74.8%;
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Pred. No. 1.2e-34
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annotation update)
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RESULT 14
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AC QAGMVI
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WA KIAUSHER R.D., Collins F.S., Wapmer L.H., Derge J.G.,

KIAUSHER R.D., Collins F.S., Wapmer L., Shemmen C.M., Schuler G.D.,

A KIAUSHER R.D., Collins F.S., Wapmer L., Shemmen C.M., Schuler G.D.,

A Altschul S.F., Zeeberg B., Buetow K.H., Schaefar C.F., Bhat N.K.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

A Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Willaho L., Bonaldo M.F., Casrain C.P., Prange C.,

A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

A Brownstein M.J., Wallano N.A., Peters G.J., Abramson R.D., Wullahy S.J.,

A Holdards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.M.,

A Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

A Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

A Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

A Korebsley R.W., Touchman J.W., Green E.D., Dickson M.C.,

A Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

A Korebs R. T. Mayres M. A.,

A Kore
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Best Local S
Matches 84
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05-UUL-2004 (TrEMBLrel. 2
05-UUL-2004 (TrEMBLrel. 2
05-UUL-2004 (TrEMBLrel. 2
Hypothetical protein.
Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; (
Mammalia; Eutheria; I
NCBI_TaxID=9606;
[1]
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EMBL; BC012876; AAH12876.1; -.
PIR; S12440; S12440.
HSSP; P01842; 1AQK.
                                                                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                  Q6GMV7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hypothetical protein.
SEQUENCE 236 AA; 24712 MW; 7EC9FB3622FED957 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jones S.J., Marra M.A.;
"Generation and initial analysis of more than 15,000 and mouse cDNA sequences.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
   SEQUENCE
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                                                        NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PDRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLGQPK 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PDRFSGSKSGTSASLAITGLQAEDEADYYCQSYDGSLSGSVFGAGTKVTVLGQPK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       73.5%;
nilarity 73.0%;
Conservative
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                                                                                                Chordata;
Primates;
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Pred. No. 4.5e-34;
9; Mismatches 21
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                                                                                                Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                          sequence update)
annotation updat
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RESULT 15
QGDHW4
ID QGDHW
AC QGDHW
AC QGDHW
AC QGDHW
DT 25-OC
DT 35-OC
DT 15-OC
DT 1
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RM MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;

RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

RA Klausner R.D., Collins F.S., Wagner L., Shemmen C.M., Schuler G.D.,

RA Altschul S.F., Zeebberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Altschul S.F., Zeebberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RA Hopkins R.F., Jordan H., Moore T., Mard S.I., Wang J., Haieh F.,

RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,

RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RA Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,

RA Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.J.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RA Richards S., Worley K.C., Hale S., Garcina A.M., Gay L.J., Hulyk S.W.,

RA Willalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,

RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,

RA Hothey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,

RA Hothey G., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,

RA Kzywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,

Dan Jones T. Marya M.A.
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Best Local S
Matches 84
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Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases.
EMBL, BC073795; AAH73795.1; -.
R InterPro; IPR003599; Ig.
R InterPro; IPR003597; Ig-like.
R InterPro; IPR003597; Ig-li.
R InterPro; IPR003597; Ig-li.
R InterPro; IPR003596; Ig-MHC.
R InterPro; IPR003596; Ig-MHC.
R InterPro; IPR003596; Ig-MHC.
R InterPro; IPR003596; Ig-V.
R Pfam; PF00647; ig; 2.
R Pfam; PF00047; ig; 2.
R SMART; SM00409; IG; 2.
R SMART; SM00406; IG-MHC; UNKNOWN_1.
R PROSITE; PS00290; IG-MHC; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                 Q6DHW4 PRELIMINARY;
Q6DHW4;
25-OCT-2004 (TrEMBLrel. 28, L
25-OCT-2004 (TrEMBLrel. 28, L
25-OCT-2004 (TrEMBLrel. 28, L
(1)
SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
TISSUE=Primary B-Cells;
MEDLINE=22388257; PubMede12477932; DOI=10.1073/pnas.242603899;
MEDLINE=22388257; PubMede12477932; DOI=10.1073/pnas.242603899;
MEDLINE=22388257; Peingold E.A., Grouse L.H., Derge J.G.,
                                                                                                                                                                                                                                   Hypothetical protein.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                             NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hypothetical
SEQUENCE 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        QSVLTQPSSVSAAPGQKYTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP
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236 AA; 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                73.5%; Score 430.5;
73.0%; Pred. No. 4.50
tive 11; Mismatches
                                                                                                                                                                                                                                                                                                                                                           Created)
Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                         Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
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RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Aleschul S.F., Zeeberg B., Buecow K.H., Schaefer C.F., Bhat N.K.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh P.,
RA Diatchenko L., Warusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Warusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Ugdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Bask S.A., Moguellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Rahá S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Mullahy S.J.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.M.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.M.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.M.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.M.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.M.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.M.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.M.,
RA Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G.,
RA Rodriguez A.C., Grimwood J., Schenta A., Worley S.M., Butterfield Y.S.,
RA Krzyminski M.I., Skalska U., Smailus D.E., Scherch A., Schein J.E.,
RA Krzyminski M.I., Skalska U., Smailus D.E., Scherch A., Schein J.E.,
RA Gorders B.M., Madan A., Young A.C., Shevchenko Y., Butterfield Y.S.,
RA Rataeley R. M., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Rataeley R.M., Touchman J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Grimwood J.W., Green E.D., Dickson M.C.,
RA Rodriguez A.C., Marza M.A.,
RA Rodriguez A.C., Grimwood J.W., Scherch A., Schein J.B.,
RA Schould J. A. Scherch M.A.,
RA Rodriguez A.C., Grimwood J.W., Green E.D.
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DRFSGSKSGTSASLAIGGLRSEDEAHYYCAAWDDSMSGPGYAFGTGTKVTVLRQPK 135
                                                                                                                                                                                                                                                              71.8%; Score 421; DE ilarity 70.7%; Pred. No. 3.9e Conservative 11; Mismatches
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1.9e-33;
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seq length:
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•; <sup>ti</sup>	409	415	415.5	415.5	417	419	419	420	429	432	434.5	435.5	436	436.5	440	442.5	4
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	S36263	LIHUVO	PH0132	LIHUMM	LIHUWA	S04525	S36068	S25743	S36274	S57428	JE0246	S51148	A44151	TTHUHA	S57408	S25754	
	Ig lambda chain V	lambda		lambda	lambda chain	Ig lambda chain pr	Ig lambda chain -	Ig lambda chain -	Ig lambda chain V	Ig light chain V-J	Ig lambda chain NI	antibody light cha	Ig lambda chain V	Ig lambda chain V-	Ig lambda chain V-	Ig lambda chain -	

## ALIGNMENTS

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RESULT 2
S47009
Ig lambda chain V1-J3 region - human
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Species: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 21-Jan-2000
C;Accession: S47009
R;Mahmoudi, M.; Gasyna, E.; Denomme, G.; Edwards, J.; Bell, D.; Cairns, E. submitted to the EMBL Data Library, July 1994
A;Bescription: The role of the immunoglobulin heavy chain in human anti-DNA
A;Reference number: S47009
A;Accession: S47009
A;Accession: S47009
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Molecule type: mRNA
A;Residues: 1-130 <KIS2>
A;Residues: 1-130 <KIS2>
A;Cross-references: RMBL:X14583
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;1-20/Domain: signal sequence #status predicted <SIG>
F;21-235/Product: Ig lambda chain #status predicted <MAT>
F;150-218/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Submitted to the EMBL Data Library, March 1989

A;Reference number: S05270

A;Accession: S05270

A;Accession: S05270

A;Rolecule type: mRNA
A;Residues: 1-235 < KIS1>
A;Cross-references: EMBL:X14583; NID:g33394; PIDN:CAA32725.1; PID:g33395

A;Kishimoto, T.; Okajima, H.; Okumoto, T.; Taniguchi, M.

Nucleic Acids Res. 17, 4385, 1989

A;Title: Nucleotide sequences of the cDNAs encoding the V-regions of H- and L-chains
A;Reference number: S04601; MUID:89296497; PMID:2500644

A;Accession: S04601
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Simi
hes 108;
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                                                                                                          the immunoglobulin heavy chain in human anti-DNA antibody bin
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A;Molecule type: mRNA
A;Residues: 1-111 <MAH>
A;Residues: 1-111 <MAH>
A;Residues: 1-111 <MAH>
C;Cross-references: EMBL: Z35495; NID:g517346; PIDN:CAA84629.1; PID:g517347
A;Cross-references: EMBL: Z35495; NID:g517346; PIDN:CAA84629.1; PID:g517347
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-91/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                               Ig lambda chain V region (Zim) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 31-Dec-1988 #sequence_revision 31-Dec-1988 #text_
C;Accession: A29700
R;Eulitz, M.; Breuer, M.; Linke, R.P.
Biol. Chem. Hoppe-Seyler 368, 863-870, 1987
A;Title: Is the formation of AL-type amyloid promoted by A,Reference number: A29700; MUID:87299022; PMID:3620114
A;Accession: A29700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Status: preliminary
A;Molecule type: mRNA
A;Molecule type: mRNA
A;Residues: 1-130 < HUG>
A;Residues: 1-130 < HUG>
A;Cross-references: GB:X52109; NID:g31454; PIDN:CAA36343.1; PID:g31455
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;34-110/Domain: immunoglobulin homology <IMM>
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A29700
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C;Date: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 21-Jan-2000
C;Accession: S09712
                                                                                A; Molecule type: protein
A; Residues: 1-113 <EUL>
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
F;14-90/Domain: immunoglobulin homology <IMM>
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A;Title: Nucleotide sequences and three-dimensional modelling
A;Reference number: S09710; MUID:90262535; PMID:2111699
A;Accession: S09712
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R;Hughes-Jones, N.C.; Bye, J.M.; Beale, Biochem. J. 268, 135-140, 1990
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Pred. No. 2.6e-39;
Score 534; DB z; r
Pred. No. 7.2e-39;
"" "" amatches 8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 538; DB 2;
Pred. No. 3.8e-39;
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RESULT 6
S24321
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
P;147-215/Domain: immunoglobulin homology <IMM>
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A;Title: V(lambda) and J(lambda)-C(lambda)
A;Reference number: S16439; MUID:91257162;
A;Accession: $25742
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C;Accession: S25742
R;Combriato, G; Klobeck, H.G.
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S25742
                                                                                                                                                                                                        A;Cross-references: EMBL:X64134; NID:g32808; PIDN:ChA45495.1; C;Superfamily: immunoglobulin V region; immunoglobulin homolog C;Keywords: heterotetramer; immunoglobulin F;34-110/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                             R;Aucouturier, P.; Khamlichi, A.A.; Preud'homme, J.L.; Bauwens, M.; Touchard, G.; Cogne, Biochem. J. 285, 149-152, 1992
A;Title: Complementary DNA sequence of human amyloidogenic immunoglobulin light-chain pra;Reference number: $24319; MUID:92344562; PMID:1379039
                                                                                                                                                                                                                                                                                                                                                                                                                   Ig lambda chain precursor - human
C;Species: Homo sapiens (man)
C;Date: 02-Dec-1993 #sequence_revision 10-Nov-1995 #text_change 21-Jan-2000
C;Accession: S24321
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A;Molecule type: mRNA
A;Residues: 1-232 <COM>
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C; Species: Homo sapiens (man)
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A;Molecule type: mRNA
A;Residues: 1-131 <AUC>
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                                                                                      1 QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP
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                                                                 OSVLTQPPSVSAAPGQKVTISCSGSSSNIANNYVSWYQQLPGAAPKLLIYENVKRPSGIP
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89.5%;
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Pred. No. 4e-38;
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Pred. No. 1.8e-38;
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PMID:1904362
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lambda chain V-I region (Nig-64) - human (tentative sequence

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A;Cross-references: GDB:119342; OMIM:147240
A;Map position: 22q11.2-22q11.2
C;Complex: An immunoricality
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A;Map position: 22q11.2-22q11.2
C;Complex: An immunoglobulin heterotetramer subunit consists of two identical light (kap hain disulfide bonds. In some cases, such as IgA and IgM, the subunits associate into la C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: blocked amino end; heterotetramer
F;15-91/Domain: immunoglobulin homology <a href="https://domain:immunoglobulin">https://domain:immunoglobulin</a> homology <a href="https://domain:immunoglobulin">https://domain:immunoglo
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P;20-130/Product: Ig lambda chain V-I region (BL2) #status predicted <MAT>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic Acids Res. 12, 8407-8414, 1984
A;Title: Molecular cloning of a human immunoglobulin lambda chain variable sequence.
A;Reference number: A01966; MUID:85062823; PMID:6095199
A;Accession: A01966
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R; Tsujimoto, Y.; Croce, C.M.
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A;Title: Comparative studies on the structure of the light chains of human immunoglobul. A;Reference number: A91970; MUID:83186114; PMID:6404900
                                                                                                                                                                                                                                                                                             F;34-110/Domain: immunoglobulin homology <IMM>F;116-130/Region: J segment
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A; Residues: 1-130 <TSU>
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C;Date: 03-Aug-1984 #sequence_revision 03-Aug-1984 #text_change 09-Jul-2004
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QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP
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                                                                               Score 523; DB 1;
Pred. No. 7.1e-38;
5; Mismatches 6
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Pred. No. 5e-38;
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A; Residues: 1-110, 'W' <JON>
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A; Residues: 1-111 < MAR>
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Best Local :
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Date: 22-Jan-1993 #sequence_revision 22-Jan-1993 #text_change 20-Jun-2000
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                                                                                                                                    1 QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP 60
                                                                                                                                                                            h 88.7%; Score 520; DB 2; Similarity 89.2%; Pred. No. 1.1e-37; 99; Conservative 3; Mismatches 9
                            DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLG
                                                                                     QSVLTQPPSVSAAPGQXVTISCSGSSSNIGNNYVSWYQHLPGTAPNLLIYDNNKRPSGIP
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DRFSGSKSGTSATLGITGLQTGDEADYYCGTWDGRLTAAVFGSGTKVTVLG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bonnert, T.P.; McCafferty, J.; Griffiths, A.D.; Winter,
                                                                                                                                                                                                                                                                                                                                                                                                                                                          October
                                                                                                                                                                                                                       Length 111
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A; Molecule: type: protein
A; Residues: 1-111 < LAN>
A; Cross-references: UNIPROT: P01701 R;Langer, B.; Steinmetz-Kayne, M.; Hilschmann, N. Hoppe-Seyler's Z. Physiol. Chem. 349, 945-951, 1968 A;Tille: The complete amino acid sequence of Bence-Jones A;Reference number: A01964; MUID:69060892; PMID:4177823 C;Species: Homo sapiens (man)
C;Date: 24-Apr-1984 #sequence\_revision 24-Apr-1984 #text\_change A, Accession: A01964 C;Accession: A01964 Ig lambda chain V-I region (New) - human
C;Species: Homo sapiens (man) RESULT 10 THUNW Comment: This a Bence Jones protein protein New (lambda type). Subg 09-Jul-2004

A; Map position: 22q11.2-22q11.2
C; Complex: An immunoglobulin herrotetramer subunit consists of two identical light (kap C; Complex: An immunoglobulin bome cases, such as IgA and IgM, the subunits associate into la L; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: blocked amino end; heterotetramer
F; 15-91/Dömain: immunoglobulin homology < IMM>
F; 1, Modified site: blocked amino end (Gln) (probably pyrrolidone carboxylic acid) #status
F; 22-89/Disulfide bonds: #status predicted

A;Cross-references: GDB:119342; OMIM:147240

C; Genetics:

;Gene: GDB:IGLV@

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RESULT 11
S47185
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C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revi
C;Accession: S36050
R;Williams, S.C.
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A;Molecule type: mRNA
A;Residues: 1-111 <MCI>
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R;MCIntosh, R.S.; Tandon, N.; Metcalfe, R.A.; Weetman, A.P. submitted to the EMBL Data Library, June 1994

A;Description: Cloning and analysis of IgM anti-thyroglobul A;Reference number: S47181

A;Accession: S47185
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C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995 #text_change 21-Jan-2000
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                                                                                                                                                                                                                                                                                           submitted to the EMBL Data Library, A;Reference number: $36046 A;Accession: $36050 A;Status: preliminary
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A; Residues: 1-98 <WIL>
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Matches 93
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                                                                       QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP
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                 DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSA 98
                                                   86.7%;
nilarity 86.5%;
Conservative
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Pred. No. 8.7e-35;
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Pred. No. 1.2e-36;
                                                                                                                     Score 485; DB 2;
Pred. No. 9.4e-35;
1; Mismatches 4
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RESULT 14
S78057
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A;Residues: 1-233 <COM>
A;Residues: 1-233 <COM>
A;Ressidues: 1-233 <COM>
A;Cross-references: EMBL:X57817; NID:g33733; PIDN:CAA40954.1; PID:g33734
A;Cross-references: EMBL:X57817; NID:g33733; PIDN:CAA40954.1; PID:g33734
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;148-216/Domain: immunoglobulin homology <IMM>
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A;Title: V(lambda) and J(lambda)-C(lambda)
A;Reference number: S16439; MUID:91257162;
A;Accession: S25752
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C;Species: Homo sapiens (man)
C;Date: 22-Nov-193 #sequence_revision
C;Accession: S25752
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: EMBL:X54438; NID:g37920; PIDN:CAA38307.1; PID:g37921 R;Harindranath, N.; Goldfarb, I.S.; Ikematsu, H.; Burastero, S.E.; Wilder, Int. Immunol. 3, 865-875, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ig lambda chain precursor V-J region (clone mAB 61VL) - human (fragment) C;Species: Homo sapiens (man) C;Dace: 19-Nov-1997 #sequence_revision 05-Dec-1997 #text_change 23-Jul-1C;Accession: 878057; 823722
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R; Combriato, G.; Klobeck, H.G.
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A;Residues: 20-130 <HAW>
A;Cross-references: EMBL:X54438
                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Reference number: S23716; MUID: 92031262; PMID: 1718404
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Reference number: 878051
A;Accession: 878057
A;Molecule type: mRNA
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F;34-110/Domain: immunoglobulin homology
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nilarity 79.8%;
Conservative :
                                                                                                                                                     81.4%; Score 477; DB 2 clarity 80.2%; Pred. No. 6e-34; Conservative 10; Mismatches
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Pred. No. 2.2e-34;
9; Mismatches 14
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PMID:1904362
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RESULT 15
A42193
19 lambda chain (BJP-DIA) - human
N;Alternate names: amyloid fibril protein AL-DIA; Bence Jones protein; Ig lambda chain (C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 03-Mar-1994 #sequence revision 07-Apr-1994 #text_change 21-Jan-2000
C;Accession: A42193; S18297
R;Klafki, H.W.; Kratzin, H.D.; Pick, A.I.; Eckart, K.; Karas, M.; Hilschmann, N.
Biochemistry 31, 3265-3272, 1992
A;Title: Complete amino acid sequence determinations demonstrate identity of the urinary A;Reference number: A42193; MUID:92207944; PMID:1554711
A;Residues: 1-216 < KLA-
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin; pyroglutamic acid
F;131-199/Domain: immunoglobulin pyroglutamic acid
F;11/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
Search completed: October 13, 2005, 03:11:47 Job time : 24.485 secs
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                                                                                                                                          61 DRFSGSKSGTSATIGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLGQPK 114
                                                                                                                61 GRESGSKSGTSASLAVSGLQSEDEADYYCATWDDSVNGWVPGGGTKLTVLGQPK 114
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Abg73422 Human IgG
Abg75348 Antibody
Aae39158 IgG3 anti
Ad809288 Human CM
Aar12263 Anti-huma
Aao31147 Human dia
Ad922405 Human ant
Adp22403 Human ant
Abg22403 Human BLy
Adg96076 Single ch
Abm8428 Human BLy
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Abp45235 Human BLy
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ALIGNMENTS

RESULT 1 ABB07187 X B X B X S X S X S X B X B 13-MAR-2002 (first entry) Neuromodulatory; central nervous system; CNS; sHIgM22; LYM 22; AKJR4; ebvHigM MsI19D10; ebv HIgM CB2bG8; CB2iE12; CB2iE7; MSI19E5; virucide; antiparkinsonian; neuroprotective; nootropic; vulnerary. sHigM22 light chain variable region clone II amino acid sequence ABB07187; ABB07187 standard; protein; 114 15-NOV-2001. WO200185797-A1 Homo sapiens.

(MAYO-) MAYO FOUND MEDICAL EDUCATION RES 10-MAY-2000; 2000US-00568351. 30-MAY-2000; 2000WO-US014902

· · · · <u>· · · ·</u>

Rodriguez M, Miller DJ,

WPI; 2002-066596/09. N-PSDB; ABA94244.

Novel neuromodulatory agent (a human IgM monoclonal antibody), promoting neurite outgrowth, regeneration, remyelination and neuroprotection in central nervous system, useful to treat post-infectious encephalomyelitis.

Claim 23; Fig 18; 219pp; English.

#X#X#X#X####XXXDDDDDDDD The invention provides a neuromodulatory agent (I) capable of promoting neurite outgrowth, regeneration, remyelination and neuroprotection in central nervous system (CNS). (I) is capable of inducing remyelination, "promoting cellular proliferation of glial cells, and promoting C2+ signaling with oligodendrocytes. An humanised antibody to (I) can be selected from antibody sHIGM2 (LYM 22), ebvHigM MSI19D10, ebv HIGM CB2D58, AKJR4, CB21E12, CB21E7 or MSI19E5. (I) is useful for stimulating remyelination of CNS axons, stimulating proliferation of glial cells in CNS axons, or treating demyelinating disease of CNS in a mammal in need of such therapy. (I) is capable of binding to structures and cells within

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               are useful for preventing infection by a bacterium, virus or like pathogen that causes demyelination or other neurodegenerative condition in a subject. Methods where (I) is administered to a pathent are useful for treating multiple sclerosis, Parkinson's disease, Alzheimer's disease, amyotrophic lateral sclerosis (ALS), a viral demyelinating disease, CNS diseases, and other conditions in the CNS where nerves are damaged as by trauma. The present sequence represents the sHigM22 light
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Neuromodulatory; central nervous system; CNS; sHIgM22; LYM 22; AKJR4; ebvHigM MsI19D10; ebv HIgM CB2bG8; CB2iE12; CB2iE7; MSI19E5; virucide; antiparkinsonian; neuroprotective; nootropic; vulnerary.
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The invention provides a neuromodulatory agent (I) capable of promoting neurite outgrowth, regeneration, remyelination and neuroprotection in central nervous system (CNS). (I) is capable of inducing remyelination, promoting cellular proliferation of glial cells, and promoting Ca2+
                                                                                                                                                                                                                                                                                          Rodriguez M, Miller DJ,
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                                                                                                                                                   Novel neuromodulatory agent (a human IgM monoclonal antibody), pronneurite outgrowth, regeneration, remyelination and neuroprotection central nervous system, useful to treat post-infectious
                                                                                                                                                                                                                                                                                                                                                                 10-MAY-2000; 2000US-00568351.
                                                                                                                                                                                                                                N-PSDB; ABA94217
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
                                                                                                                                                                                                                                                  2002-066596/09.
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RESULT 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches 112;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                                                         29-APR-1994;
08-AUG-1996;
07-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                         demyelination disease; multiple scierosis; central raxon; glial cell proliferation; Theiler's murine encephalomyelitis virus infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 114 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                              Human; antibody; IgM; remyelination; neuronal growth; autoantibody; demyelination disease; multiple sclerosis; central nervous system;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human anti IgM antibody sHIgM22 VL protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADI26656;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADI26656 standard; protein; 114
WPI; 2004-119219/12
                                 Rodriguez M,
                                                                                                                                                                                                                                     13-NOV-2001;
                                                                                                                                                                                                                                                                      02-OCT-2003
                                                                                                                                                                                                                                                                                                        US2003185827-A1
                                                                                                                                                                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                            spinal cord injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-APR-2004 (first entry
                                                                                                                          30-MAY-2000;
                                                                                                                                              28-MAY-1999;
                                                                      (MAYO-) MAYO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ocal Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     QSVLTQPPSVSAAPGQKVT1SCSGSSSN1GNNFVSWYQQLPGTAPRLL1YD1TKRPSG1P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DRFSGSKSGTSATLGITGLQTGDEADYYCGTWDSSLSAVVFGGGTKLTVLGQPK 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                        FOUND.
                                                                                                       99US-00322862.
2000US-00580787.
2000US-00730473.
                                                                                                                                                                                                                                     2001US-00010729.
                                     Miller
                                                                                                                                                           94US-00236520.
96US-00692084.
97US-00779784.
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                                   Ŋ,
                                     Pease LR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 582; DB 5;
Pred. No. 8.8e-41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                  CNS injury
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gape
                                                                                                                                                                                                                                                                                                                                                                                                                                      CNS;
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17

Human diagnostic and therapeutic pprotein SEQ ID NO:4542

New human immunoglobulin M antibody for treating or preventing demyelinating disease of the central nervous system in a human domestic animal, such as multiple sclerosis. animal, such as multiple sclerosis. or a

Claim 6; Fig 36; 159pp; English.

The invention relates to an antibody (I) produced by injecting an antibody, where the peptide comprises a human anti-1gM antibody fragment CC given in the specification, or active fragments. Also included are commandal (comprising administering a monoclonal antibody, or maxtures, monomers, active fragments, or recombinant antibodies derived from it, characterised by their ability to bind structures and cells within the CC (CNS, including oligodendrocytes), stimulating dhe proliferation of glial CC cells in CNS axons in a mammal (comprising administering a monoclonal antibody, or mixtures, monomers, active fragments, or recombinant antibodies derived from it, characterised by their ability to bind structures and cells within the CNS), treating a monoclonal antibody, or mixtures, monomers, active fragments, or recombinant compositional antibody, or mixtures, monomers, active fragments, or recombinant antibodies derived from it, characterised by their ability to bind structures and cells within the CNS), and to stimulate remyelination of demyelinating attribudy, or mixtures, monomers, active fragments, or recombinant antibody, or mixtures, monomers, active fragments, or composition of the CNS), stimulating remyelination of CNS axons in a mammal (comprising administering a compositional antibody, or mixtures, monomers, active fragments or compositional antibody, or mixtures, monomers, active fragments of the cNS axons in a mammal (compositional antibody), an asset of the cNS, and to stimulate remyelination of GNS axons in a mixture, the proliferation of Spial antibody, an antibody and antibody

Sequence 114

Matches

114;

Query Match Best Local Similarity

Length 114; Indels

S 밁 S 61 61 Ъ -QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLGQPK 114 DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLGQPK 114 QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP 99.3%; Score 582; DB 8; ilarity 100.0%; Pred. No. 8.8e-41; Conservative 0; Mismatches 0; 0 Gaps 60 60 0

RESULT 4 ABM84293 SPASSE ABM84293; ABM84293 standard; protein; 234

18-NOV-2004

(first entry)

Mooney EM, Delegeane AM, Panesar IS, Banville SC, Reddy TP; Stevens KA, Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin Peralta CH, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL; Lagace RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton Xu Y, Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, G Patury S, Shi X, Suarez CJ; gene therapy; human diagnostic and therapeutic polynucleotide; dithp WPI; 2004-329368/30. Schmidt JP, 12-SEP-2002; 12-SEP-2002; 12-SEP-2003; 2003WO-US028227 WO2004023973-A2 Homo sapiens Harthshorne TA, (INCY-) INCYTE CORP Wright RJ, 2002US-0410259P. 2002US-0410260P. ht RJ, Bruns C Suchorolski MT, a CM, mary... MT, Altus CM, Banv Marjanovic MM, Pitts SJ, Shen F; Reddy TP; .P, Gerstin EH; Elder LV; Kirton ES; Gietzen D;

in gene mapping. New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy or

Claim 27; Page; 190pp; English

The invention relates to novel diagnostic and therapeutic polynucleotides conserved from one of the 2722 sequences defined in the specification. A CC polynucleotide of the invention may have a use in gene therapy. The human CC diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be used to diagnose a particular condition, disease or disorder associated CC with human molecules, e.g. cell proliferative disorder, endocrine autoimmune/inflammatory disorders, developmental disorders, or clinections caused by virus, bacteria, fungi or parasite. The dithp cm collecules may also be used in genetic mapping, in identifying individuals cc from minute biological samples, in detecting single nucleotide polymorphisms, as molecular weight markers, and for somatic or germline cc polymorphisms, as molecular weight markers, and for somatic or germline cc invention. Note: The sequence represents a dithp protein of the constitution. The present sequence represents is not represented in the printed specification, but was obtained in electronic format directly from WIPO at www.wipo.int/pct/en/sequences/listing.htm

Query Match Best Local ( Matches 93.4%; Similarity 93.9%; Conservative Score 547.5; DB ( Pred. No. 1.3e-37

Sequence

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20 1 QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNYVSWYQQLPGTAPKLLIYDNNKRPSGIP

Mismatches

DB 8;

Length Indels

234;

1;

Gaps

60 79

61 DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLGQPK 114 DRFSGSKSGTSATLGITGLQTGDBADYYCGTWDSSL-MVVFGGGTKLTVLGQPK 132

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ABG73422 RESULT: 5 ABG73422 standard; protein; 234 AA

AC ABG73422;

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RESULT 6
ABG75338
ID ABG7
XX
AC ABG7
AC ABG7
XX
AC ABG7
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                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a reagent comprising an IgG anti-Rh antibody chaving two lambda light chains, two gamma 3 heavy chains and a polymeric tailpiece attached to the carboxy terminal end of the heavy chain. The reagent is useful in a haemagglutinin test to determine the presence of one or more Rh (rhesus) antigens on red blood cells. The reagent is also useful for testing red blood cells to determine the presence of one or more Rh factors, by contacting red blood cells with the reagent for a sufficient time and at a sufficient temperature to cause agglutination of red blood cells which have one or more Rh factors present, and determining whether the red blood cells agglutinate to determine the presence of one or more Rh factors present, and accurate Rh phenotyping of blood and is used in the one-step haemagglutinin method. This sequence represents a human IgG anti-Rh antibody lambda collight chain polypeptide
                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local
                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Reagent for identifying rhesus antigen on red blood cells, comprises antibody hybrid having IgG anti-Rh antibody which has polymeric tail piece attached to carboxy terminal end of each of IgG antibody heavy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     IgG anti-rhesus antibody; Rh; gamma 3 heavy chain; lambda light chain; haemagglutinin; rhesus antigen; red blood cell; agglutination; antibody; Rh phenotyping; human.
                                                                                                                ABG75338 standard;
                                                                                                                                                                                                                                                                                                                                                                                             Sequence 234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Col 13-16; 14pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Morrison SL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          11-AUG-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-AUG-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24-APR-2003 (first entry
               Antibody single chain framework polypeptide SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (REGC ) UNIV CALIFORNIA.
                                                 22-APR-2004
                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2003-208833/20
                                                                                                                                                                                                                                                                                                                              103;
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                                                                                                                                                                                                                  DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLGQPK 114
                                                                                                                                                                                                                                                             QSVLTQPPSVSAAPGQKYTISCSGSSSNIGNNYVSWYQHFPGTAFKLLIYDNNQRPSGIP
                                                                                                                                                                                                                                                                                      QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP
                                                                                                                                                                                                DRFSGSKSGTSATLGITGLQTGDEADYYCGTWDSSLSAVLFGGGTEVTVLGQPK 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       anti-rhesus antibody lambda light chain polypeptide
                                                                                                                                                                                                                                                                                                                              Conservative
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                                                 (first entry)
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                                                                                                              protein; 111
                                                                                                                                                                                                                                                                                                                                           92.8%;
90.4%;
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                                                                                                                                                                                                                                                                                                                                             Score 544; DB 6;
Pred. No. 2.5e-37;
                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                 8
                                                                                                                                                                                                                                                                                                                              6;
                                                                                                                                                                                                                                                                                                                                                              Length 234;
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RESULT 7
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UX 2 X E X E X X X X E F F F X
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Best Local Simi
Matches 104;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to a single chain antibody framework. The single chain framework, antibody or antibody fragment, is useful in target validation, diagnostic applications, library construction or therapeutic applications. The framework sequences are used in the identification of a conserved framework residue class selected from polar but uncharged R groups, positively charged R groups, negatively charged R groups, negatively charged R groups, negatively charged R groups, and special amino acids. The framework sequences may also be used in the identification of at least one conserved framework sequence, where the conserved framework sequence is 2 -5, 5-10 or 10-25 residues, and where the conserved framework sequence has gaps. The nucleic acid molecule is used in gene therapy. The present sequence is a polypeptide of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 111 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New single chain immunoglobulin frameworks that demonstrate enhanced stability in the intracellular environment, useful for target validates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-022852/02.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-MAY-2002; 2002US-0382649P.
03-JAN-2003; 2003US-0438256P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antibody; framework; stability; intracellular;
                                                                                                  Rh phenotyping; prophylactic.
                                                                                                                    Anti-rhesus antibody; Rh; immunoglobulin G; IgG; haemagglutinin test Rh phenotyping; therapeutic; haemolytic disease; Rh incompatibility;
                                                                                                                                                                                             18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-MAY-2003; 2003WO-EP005324
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO2003097697-A2
                            Misc-difference
                                                                        Unidentified
                                                                                                                                                              IgG3 antibody lambda light chain protein
                                                                                                                                                                                                                                                      AAE39158 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ESBA-) ESBATECH AG
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                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       construction,
                                                                                                                                                                                                                                                                                                                                               DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLG 111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Page 48; Opp; English.
                                                                                                                                                                                                                                                                                                                                 DRFSGSKSGTSATLGITGLQTGDBADYYCGTWDSSLSGVVFGGGTKLTVLG 111
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                            (first entry)
                          Location/Qualifiers
                                                                                                                  therapeutic; haemolytic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  92.7%;
93.7%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 543; DB 8;
Pred. No. 1.5e-37;
3; Mismatches 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          or diagnostic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 111;
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/note= "Encoded by GAT"

US2003143643-A1

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ADS09284
ADS09284
ADS09284
ADS09284
AC ADS0
XX ADS0
XX ADS0
XX ADS0
DT 18-N
XX C-Me
KW OPH
KW OPH
KW OPH
KW OPH
KW OPH
KW PBOT
XX PBOT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 92.0%;
Best Local Similarity 89.5%;
Matches 102; Conservative
                                                                                                                                                                                                                                                                                                                                                       c-Met; tyrosine kinase antibody; antigen binding; cytostatic; ophthalmological; antiinflammatory; analgesic; vasotropic; antipsoriatic; osteopathic; cancer; tumour; ophthalmic disease; glaucoma; retinitis; retinopathy; uveitis; ocular photophobia; macular degeneration; pain; acute injury; eye; hyperproliferative disorder; restenosis; angioplasty; psoriasis; HGF; osteoporosis; cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to hybrid anti-Rhesus (Rh) antibody comprising an immunoglobulin (Ig)G anti-Rh antibody which comprises two light chains and two heavy chains where the heavy chains have a carboxy terminal with a polymeric tailpiece and an amino terminal. The antibodies are used in a one-step haemagglutinin test to provide Rh phenotyping of blood and as therapeutic or prophylactic agents for preventing haemolytic disease in newborns due to Rh incompatibility. The present sequence is IgG3 antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hybrid immunoglobulin G anti-Rhesus antibody for use in one-step kh phenotyping of blood or as a prophylactic agent to prevent hemolytic disease in newborns due to Rh incompatibility, comprises a polymeric
                              13-FEB-2003; 2003US-0447073P
                                                                                                                                                                                                                                                                                                Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human c-Met protein tyrosine kinase antibody, PGIA-4-A3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-NOV-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADS09288 standard; protein; 248
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                                                                                            11-FEB-2004; 2004WO-IB000503
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A. 198

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61 DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLG 111

DRFSGSKSGTSATLGITGLQTGDEADYYCGTWDSSLSGVVFGGGTKLTVLG

248

QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP 60

Query Match 92.9
Best Local Similarity 92.9
Matches 103; Conservative

92.0%;

Score 539; DB 8; Length 248; Pred. No. 7e-37; 3; Mismatches 5; Indels

<u>..</u>

Gaps

Sequence 248 AA;

RESULT 9
AAR12263
ID AAR12
XX
AC AAR1
XX
DT 25-M
DT 15-A
DT 15-A
XX
XX
XX
XX
XX
MON
DR
XX
MON
ON
MW haem
XX
ON
ON
HOMO

25-MAR-2003 15-AUG-1991

(revised) (first entry)

AAR12263;

AAR12263 standard; protein;

Monoclonal haemolytic

antibody; rhesus D; blood-typing; CDR, disease of the newborn; HDN.

Anti-human

RhD FOG-B

MAb (VL chain).

Homo sapiens

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The invention relates to a novel c-Met protein tyrosine kinase antibody cor its antigen binding portion that specifically binds to c-Met. The c-Met antibody comprises any one of 160 fully defined sequence of 238, CC 244, 240, 250, 251, 242, 245, 247, 246, 253, 249, 243, 241, etc., amino acids as given in the specification, or its fragment. The invention CC further comprises: a pharmaceutical composition comprising the c-Met protein tyrosine kinase antibody and a carrier; an isolated cell that comprises and a carrier; an isolated cell that produces the c-Met protein tyrosine kinase antibody; and an isolated comprises and uncleic acid sequence that encodes a heavy chain or its antigen-binding portion or light chain or its compitation.

The c-Met protein tyrosine kinase antibody has cytostatic, and osteopathic activities. The c-Met protein tyrosine kinase antibody is useful for the manufacture of medicament for the treatment of cancer or composing the presence or ligation of c-Met expressing tissue. The c-Met protein tyrosine kinase antibody is useful for the treatment or prevention of ophthalmic composing the presence or ligation of c-Met expressing tissue. The c-Met protein tyrosine kinase antibody is useful for detecting c-Met in a biological sample in vitro or in vivo. The c-Met protein tyrosine kinase antibody is useful in the treatment or prevention of ophthalmic composition of c-Met protein tyrosine kinase antibody is useful for detecting c-Met in a biological sample in vitro or in vivo. The c-Met protein tyrosine kinase antibody is useful for detecting c-Met in a composition of c-Met protein tyrosine kinase antibody is useful for the treatment of associated with acute injury to the cycle of the treatment of composition of composition of composition of composition of composition of
                                                           sequence represents the protein of a \bar{\mbox{\footnotesize phage}} display generated human c-Met antibody of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel c-Met protein tyrosine kinase antibody or its antigen-binding portion specifically binding to c-Met, useful for manufacture of medicament for treating cancer or tumor and for treatment of ophtha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2004-616044/59.
N-PSDB; ADS09348.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      diseases such as glaucoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (PHAA ) PHARMACIA CORP.
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Matches 103;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The DNA sequence of eleven monoclonal antibodies are represented in Q119145-57. Synthetic genes, for both heavy and light chains may be created by combining selected CDR 1, 2, and 3 regions, which may be selected from different antibody mols. having varied binding specificity. The chimaeric anti-RhD antibodies can be used for diagnosis and therapy, and are capable of providing blood-typing reagents of high specificity and reliability. They can also be used in passive immunisation to prevent haemolytic disease of the newborn. (Updated on 25-MAR-2003 to correct PA field.) (Updated on 25-MAR-2003 to correct PI field.)
          Human; protein coordinate data; heavy chain variable domain; VH; cancer; complementarity determining region; CDR; flight chain variable domain; VL; TRAIL receptor 7; TR7; tumour necrosis factor; KILIER; death receptor 5; DR5; TRAIL receptor 2; TRAIL-R2; TNF-related apoptosis-inducing ligand; Kaposis sarcoma; central nervous system; medulloblastoma; neuroblastoma; glioblastoma; central nervous system; medulloblastoma; neuropic; glioblastoma; gratt versus host disease; antibody therapy; nootropic; AIDS; acquired immune deficiency syndrome; neurodegenerative disorder;
                                                                                                                                  Human CM085Cl1 scFv protein that specifically binds TR7
                                                                                                                                                                 06-OCT-2003
                                                                                                                                                                                                                             AAO31147 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 111 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Fig 2; 32pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DNA encoding complementary determining regions - of human antibodies, useful in prodn. of monoclonal antibodies and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1991-178104/24.
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                                                                                                                                                                                                                                                                                                                      DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVL 110
                                                                                                                                                                                                                                                                                                                                                                 QSVLTQPPSVSAAPGQKVTISCSGTSSNIGNNYVSWYQQLPGTAPKLLIYDNNKRPSGIP
                                                                                                                                                                                                                                                                                                     DRFSGSKSGTSATLGITGLRTGDEADYYCATWDSSLSAVVFGGGTKLTVL 110
                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                (first entry)
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neuroprotective; antibody
                                                                                                                                                                                                                             protein;
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Pred. No. 3.
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for passive
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RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                          The invention relates to an isolated antibody or its fragments such as CVHCDR1 (heavy chain variable domain complementarity determining region), VCCWR1 (light chain variable domain complementarity determining region), VLCDR1 (light chain variable domain complementarity CVHCDR2, VHCDR3, VLCDR1 (light chain variable domain complementarity CVHCDR2), VLCDR2 or VLCDR3. The antibody or its fragment CVHCDR3 is factor; TNP-related CVHCDR3, TR2 is also referred to as CVHCDR3, receptor 2 (TRAIL receptor 7 (TR7), TR7 is also referred to as CVHCDR3, receptor 2 (TRAIL-R2), death receptor 5 (DR5) and KILLER. The CVHCDR3 or Its fragment is useful for treating, preventing or CVHCDR3 or Its fragment is useful for treating, preventing or CVHCDR3 or CVHCDR3, The CVHCDR3 or CVHCDR3, THE CVHCDR3 or CVHCDR3 or CVHCDR3, THE STATE ALDS (acquired immune deficiency syndrome) or CVHCDR3, The present sequence is human scFv protein that specifically chicagon. The present sequence is human scFv protein that specifically
                                                                                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                                           Query Match
Best Local
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05-APR-2002;
04-JUN-2002;
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15-AUG-2002;
13-NOV-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New antibody or its fragment, useful for treating, preventing or ameliorating a cancer, e.g. colon, breast, uterine, pancreatic, lugastrointestinal cancer, or Kaposi's sarcoma or, graft versus host
                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 2; Page 288; 301pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                N-PSDB;
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                                                                                                                           ABM84291
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                                                                                                                                                                                                                                                      133
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                                                                                                                                                                                                                DRFSGSKSGTSATIGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLG
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                                                                                                                          standard; protein; 234 AA
                                                                                                                                                                                                 DRFSGSKSGTSATLGITGLQTGDEADYYCGTWDSSLSALVFGGGTKVTVLG
                                                                                                                                                                                                                                                      QSVLTQPPSVSAAPGQKVTISCSGSTSNIGNNYVSWYQQVPGTAPKLLIYDNNKRPSGIP
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                                                                                                                                                                                                                                                                                                              Conservative
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2002US-0403370P.
                                                                                                                                                                                                                                                                                                                                                                      A,
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2002US-0384828P.
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                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                           91.8%;
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Pred. No. 8.3e-37;
5; Mismatches 4
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                                                                                                                                                                                                                                                                                                                                         Length 243
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                                                                                                                                                                                                                                                                                                              Gaps
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gene therapy; human diagnostic

and therapeutic polynucleotide; dithp

Human diagnostic and therapeutic pprotein SEQ ID NO:4540

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RESULT 12
ADP22405
ID ADP22
XX
AC ADP22
XX
AC ADP22
XX
DT 09-SE
XX
DE Human
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                                                                                                                                                                                                                                                                                                    밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The invention relates to novel diagnostic and therapeutic polynucleotides CC selected from one of the 2722 sequences defined in the specification. A CC polynucleotide of the invention may have a use in gene therapy. The human CC diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be CC used to diagnose a particular condition, disease or disorder associated CC with human molecules, e.g. cell proliferative disorders, cancerine CC disorder, neurological disorders, developmental disorders, or CC infections caused by virus, bacteria, fungi or parasite. The dithp CC molecules may also be used in genetic mapping, in identifying individuals CC from minute biological samples, in detecting single nucleotide CC polymorphisms, as molecular weight markers, and for somatic or germline CC gene therapy. The present sequence represents a dithp protein of the invention. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from without in the province of the printed specification, but was obtained in electronic format directly
                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local S
Matches 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Schmidt JP, Wright RJ, Bruns CM, Marjanovic MM, Shen F;
Harthshorne TA, Suchorolski MT, Altus CM, Pitts SJ, Elder LV;
Mooney EM, Delegeane AM, Panesar IS, Banville SC, Reddy TP;
Stevens KA, Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin EH;
Peralta CH, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL;
Lagace RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton ES;
Xu Y, Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, Gietzen D;
Patury S, Shi X, Suarez CJ;
                                       09-SEP-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy or
                                                                               ADP22405
                                                                                                                  ADP22405 standard; protein; 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 234 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 27; Page; 190pp; English.
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12-SEP-2002; 2002US-0410260P
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                                                                                                                                                                                                                                                                                                                                                                                                    local Similarity
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                                                                                                                                                                                                                                                                                             QSVLTQPESVSAAPGQKVTISCSGSSSNIGNNYVSWYQQLPGTAPKLLIYDNNKRPSGIP
                                                                                                                                                                                                                                                                                                                    QSVLTQPPSVSAAPGQXVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITXRPSGIP 60
                                                                                                                                                                                                                  DRFSGSKSGTSATLGITGLQTGDEADYYCGTWDSSL-MLGFGGGTKLTVLGQPK 132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        at
                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               www.wipo.int/pct/en/sequences/listing.htm
                                                                                                                                                                                                                                                                                                                                                                                                  91.7%;
92.1%;
                                                                                                                                                                                                                                                                                                                                                                           Score 537.5; DB 8;
Pred. No. 8.8e-37;
2; Mismatches 6;
                                                                                                                                                                                                                                                                                                                                                                                                                    Length 234;
                                                                                                                                                                                                                                                                                                                                                                             1;
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Human anti-TNFa antibody light chain variable region SEQ ID NO:311.

Query Match

91.6%;

Score 537;

DB 8 ;

Length 110;

Sequence 110 AA;

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The present invention describes a human monoclonal antibody (I) that CC specifically binds to tumour necrosis factor-alpha (TNFa) and comprises: CC (a) a heavy chain complementarity determining region 1 (CDR1) having the CC two.fully defined 5 amino acid sequence (31, ADP2241) or (32, ADP22421); CC and (b) a light chain CDR1 having the two fully defined 11 amino acid compenses (33, ADP22418) or (54, ADP22424). Also described: (1) assaying CC (M1) the level of TNFa in a patient sample, comprising contacting with CC (I), and detecting the level of binding between the antibody and TNFa in the sample; (2) a composition comprising the antibody or its functional conspiration of the sample; (2) a composition comprising the antibody or its functional composition comprising the antibody or its functional composition of (11); and treating (M2) an animal suffering from a composition of (11); and (11) and (11) and (12) an animal suffering the human conclonal antibody of (11); and (12) an animal in need of treatment for TNFa induced apoptosis by administering the human monoclonal antibody of (11). (1) has anabolic, antiarteriosclerotic, antiarthritic, antiarthritic, antiarteriosclerotic, antiarthritic, and and antiapoptoric antiarthritic, antiarthritic, and and antiapoptoric antiarthritic, antiarthritic, and antiarthritic, and antiarthritic, and antiarthritic, and antiarthritic, and antiarth
                                                                            disease, graft-host reactions, septic shock, cachexia, anorexia, and multiple sclerosis. The present sequence represents a human anti-TNFa antibody light chain variable region, which is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              human; monoclonal antibody; tumour necrosis factor-alpha; TNFa; anti-TNFa antibody; anabolic; antiarteriosclerotic; antiarthritic; antibacterial; antiinflammatory; antiporiatic; antirheumatic; eating-disorder; immunomodulator; immunosuppressive; nephrotropic; neuroprotective; vasotropic; antiapoptotic; TNFa antiagonist; TNF induced apoptosis; neoplastic disease; breast cancer; ovarian cancer; bladder cancer; lung cancer; glioblastoma; stomach cancer; endometrial cancer; kidney cancer; colon cancer; parostrate cancer; immuno-mediated inflammatory disease; psoriasis; restenosis; autoimmune disease; Crohn's disease; psoriasis; restenosis; autoimmune disease; Crohn's disease; psoriasis; cancer; autoimmune disease; Crohn's disease; psoriasis; cancer; autoimmune disease; Crohn's disease; psoriasis; cancer; cancer; autoimmune disease; Crohn's disease; cancer; cance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Babcook JS, Kang JS, Foord O, Green L, Peng X, Klakamp
Haak-Frendscho M, Rathanaswami P, Pigott C, Liang ML, Le
Manchulencho K, Paggioni R, Senaldi G, Qiaojuan JS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New recombinant human monoclonal antibody that specifically binds to 
Tumor Necrosis Factor-alpha, useful for treating neoplastic disease such 
as cancers, or immuno-mediated inflammatory diseases such as rheumatoid 
arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 10; SEQ ID NO 311; 213pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 02-DEC-2002; 2002US-0430729P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-DEC-2003; 2003WO-US038281
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          septic shock; cachexia; anorexia; multiple sclerosis.
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RESULT 13
ADP22403
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
(a) a heavy chain complementarity determining region 1 (CDR1) having the two fully defined 5 amino acid sequence (S1, ADP22417) or (S2, ADP22421); and (b) a light chain CDR1 having the two fully defined 11 amino acid sequence (S3, ADP22418) or (S4, ADP22424). Also described: (1) assaying (M1) the level of TNFa in a patient sample, comprising contacting with (I), and detecting the level of binding between the antibody and TNFa in the sample; (2) a composition comprising the antibody and TNFa in the sample; (2) a composition comprising the antibody or its functional fragment and a carrier; (3) treating (M2) an animal suffering from a nanimal in need of treatment for the disease by administering the human monoclonal antibody of (I); and (4) inhibiting (M3) TNFa induced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        anti-TNFa antibody; anabolic; antiarteriosclerotic; antiarthritic; antiarthribecterial; antiinflammatory; antipsoriatic; antirheumatic; eating-disorder; immunomodulator; immunosuppressive; neptrotropic; neuroprotective; vasotropic; antiapoptotic; TNFa antagonist; TNF induced apoptosis; neoplastic disease; breast cancer; ovarian cancer; bladder cancer; lung cancer; glioblastoma; stomach cancer; endometrial cancer; kidney cancer; colon cancer; pancreatic cancer; prostrate cancer; immuno-mediated inflammatory disease; prostrate cancer; immuno-mediated inflammatory disease;
                                                                                                                                                                                                                                                                                                                                             New recombinant human monoclonal antibody that specifically binds to Tumor Necrosis Factor-alpha, useful for treating neoplastic disease such as cancers, or immuno-mediated inflammatory diseases such as rheumatoid
                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-480601/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Manchulencho K, Faggioni R, Senaldi G, Qiaojuan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Babcook JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          02-DEC-2002; 2002US-0430729P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       septic shock; cachexia; anorexia; multiple sclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      rheumatoid arthritis; glomerulonephritis; atherosclerosis; psoriasis;
restenosis; autoimmune disease; Crohn's disease; graft-host reaction;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human anti-TNFa antibody light chain variable region SEQ ID NO:309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADP22403
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADP22403 standard; protein; 110
                                                                                                                                                                                                                                                                                      Example 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Haak-Frendscho
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (ABGE-) ABGENIX INC.
                                                                                                                                                                                                                                            present invention describes a human monoclonal antibody (I) that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           104;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   monoclonal antibody; tumour necrosis factor-alpha; TNFa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVL 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OSVLTOPPSVSAAPGOKVTISCSGSSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNYVSWYQQLPGTAPKLLIYDNNKRPSGIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DRFSGSKSGTSATLGITGLQTGDEADYYCGTWDSSLSAGVFGGGTKLTVL 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                      SEQ ID NO 309; 213pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kang JS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Rathanaswami P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Foord O,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred. No. 4.6e-37;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Green L,
P, Pigott
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Feng X, Klakamp
C, Liang ML, Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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RESULT 14
ABP45249
ID ABP45:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CC apoptosis in an animal by selecting an animal in need of treatment for TNFa induced apoptosis by administering the human monoclonal antibody of CC (I). (I) has anabolic, antiarteriosclerotic, antiarthritic, CC antibacterial, antiinflammatory, antipsoriatic, antirheumatic, eating-CC disorders, immunomodulator, immunosuppressive, nephrotropic, CC neuroprotective, vasotropic and antiapoptocic activities, and can be used CC neuroprotective, vasotropic and antiapoptocis, neophrastic of sease such as TNFa antagonist. The antibody (I) is useful in the preparation of CC medicament for treating TNF induced apoptosis, neophastic disease such as the stomach cancer, ovarian cancer, bladder cancer, lung cancer, glioblastoma, CC stomach cancer, endometrial cancer, kidney cancer, colon cancer, CC pancreatic cancer, and prostrate cancer; or immuno-mediated inflammatory CC diseases such as rheumatoid arthritis, glomerulonephritis, color cancer, and prostrate cancer, accompany and the sease, graft-host reactions, septic shock, cachexia, anorexia, and CC multiple sclerosis. The present sequence represents a human anti-TNFa antibody light chain variable region, which is used in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                   BLys; B lymphocyte stimulator; TNF superfamily; human; cytostatic; tumour necrosis factor; B cell proliferation; B cell differentiation; immunosuppressive; immunostimulant; immunomodulatory; antirheumatic; antiAIDS; vaccine; cancer; immune; autoimmune disorder; immunodeficiency; systemic lupus erythematosus; rheumatoid arthritis; CVID; AIDS; common variable immunodeficiency; acquired immunodeficiency syndrome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 110 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABP45249 standard; protein; 258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human BLyS binding scrv SEQ ID 1260.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-AUG-2002 (first entry)
                                                                                                                                                                                          16-MAR-2001;
21-MAR-2001;
                                                                                                                                                                                                                            16-JUN-2000;
17-OCT-2000;
                                                                                                                                                                                                                                                                               15-JUN-2001; 2001WO-US019110.
                                                                                                                                                                                                                                                                                                                  10-JAN-2002.
                                                                                                                                                                                                                                                                                                                                                     WO200202641-A1
                                                                                                                                                                                                                                                                                                                                                                                    Homo, sapiens.
                                                                                   MS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVL 110
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                                                                                                                    HUMAN GENOME SCI INC.
CAMBRIDGE ANTIBODY TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DRFSGSKSGTSATLGITGLQTGDEADYYCGTWDSSLSAGVFGGGTKLTVL 110
                                                                                     Barash
                                                                                                                                                                       ; 2000US-0212210P.
; 2000US-0240816P.
; 2001US-0276248P.
; 2001US-0277379P.
; 2001US-0293499P.
                                                                                     SC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      91.6%;
                                                                                     Chot
                                                                                     9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
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                                                                                     Vaughan
                                                                                     'n,
                                                                                       Hilbert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 110
                                                                                       Ö
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Antibodies against B Lymphocyte Stimulating polypeptides, diagnosis and treatment of cancers and immune disorders.

useful for

WPI; 2002-114799/15

Claim 1; Page 1906-1907; 3148pp; English.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cytostatic, immunosuppressive, immunostimulant, immunomodulatory, antirheumatic and antiAIDS activity and can be used in vaccines to inhibit the expression and activity of BLyS. The antibodies bind to BLyS and so may be used to detect and quantitate the presence of BLyS in biological samples and may be used in this way to diagnose disease associated with aberrant expression of BLyS. They may also be administered to treat diseases associated with aberrant BLyS expression and activity such as cancer, immune, and autoimmune disorders and diseases, e.g. systemic lupus erythematosus, rheumatoid arthritis, immunodeficiency (e.g. common variable immunodeficiency (CVID) and acquired immunodeficiency syndrome (AIDS)). ABP43990-ABP47228 represent the antibodies and fragments of the antibodies described in the method of
            Novel antibody that immunospecifically binds to a B lymphocyte stimulator (BLys), useful for detecting and treating diseases or disorders e.g. rheumatoid arthritis, asthma and leukemia.
                                                                                                                                                                                                             16-NOV-2001; 2001US-0331469P.
19-DEC-2001; 2001US-0340817P.
                                                                                                                                                                                                                                                                                                                10-JUL-2003
                                                                                                                                                                                                                                                                                                                                                     WO2003055979-A2
                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                            B cell proliferation; differentiation; scFv; myasthenia gravis; multiple sclerosis; asthma; rheumatoid arthritis; AIDS; leukaemia; carcinoma; lymphoma; antirheumatic; antiarthritic; neuroprotective; antiinflammatory; antiasthmatic; antiallergic; cytostatic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          antibody; B lymphocyte stimulator; BLyS; tumour necrosis factor;
B cell_proliferation; differentiation; scFv; myasthenia gravis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Single chain antibody that immunospecifically binds BLyS SeqID 1260.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             11-MAR-2004 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADG96076 standard; protein; 258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 258 AA;
                                                                                           WPI; 2003-505530/47
                                                                                                                                                                       (HUMA-) HUMAN GENOME SCI INC.
                                                                                                                                                                                                                                                                      14-NOV-2002; 2002WO-US036496.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   B Lymphocyte Stimulator (BLyS) polypeptides. BLyS is a member of the tumour necrosis factor (TNF) super family and induces B cell proliferation and differentiation. The antibodies of the invention have
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes novel antibodies that immunospecifically bind to B Lymphocyte Stimulator (BLyS) polypeptides. BLyS is a member of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               208
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNFVSWYQQLPGTAPXLLIYDITKRPSGIP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DRFSGSKSGTSATLGITGLQTGDEADYYCGAWDSSLSAYVFGSGTKLTVLG 258
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNYVSWYQQLPGTAPKLLIYDNNKRPSGIP 207
                                                                                                                                Barash SC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       91.1%; Score 534; DB 5; Length 258; 92.8%; Pred. No. 1.9e-36;
                                                                                                                                Choi GH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1,
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                                                                                                                                Vaughan TJ, Hilbert
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                                       This invention relates to novel antibodies that immunospecifically bind CC to B lymphocyte stimulator (BLyS). The BLyS gene has been mapped to CC chromosome 13934 and encodes a protein that is a member of the tumour necrosis factor superfamily and induces both in vivo and in vitro B cell CC chain antibody molecules (scFws) derived, preferably, it refers to single CC chain antibody molecules (scFws) derived, preferably, from the variable cheavy CPR3 region that immunospecifically bind to a polypeptide, or CC fragment invention refers to the use of such antibodies in various methods compressed invention refers to the use of such antibodies in various methods compressed invention or inappropriate function of BLyS or its receptor. As such, these compositions are useful for identifying immune disorders compositions are useful for identifying immune disorders such these compositions are useful for identifying immune disorders compositions are useful for identifying immune disorders such as AIDS and proliferative disorders including leukaemia, carcinoma and CC lymphoma. Accordingly, they can be described as exhibiting various activities such as antixheumatic, antialtergic and cytostatic. This complete sequence is a single chain antibody that binds BLyS of the printed specification, but was obtained in electronic format condirectly from WIPO at ftp. wipo.int/pub/published pct_sequences.
Sequence 258 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 1; SEQ ID NO 1260; 394pp; English.
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'Query Match 'Best Local Similarity

91.1%;

Score 534; DB 7; Length 258; Pred. No. 1.9e-36;

₽. 묽 S Ş Matches 103; 148 QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNYVSWYQQLPGTAPKLLIYDNNKRPSGIP 208 61 1 QSVLTQPPSVSAAPGQKVTISCSGSSSNIGNNEVSWYQQLPGTAPXLLIYDITKRPSGIP DRFSGSKSGTSATLGITGLQTGDEADYYCXTWDSSLSAVVFGGGTKLTVLG DRFSGSKSGTSATLGITGLQTGDEADYYCGAWDSSLSAYVFGSGTKLTVLG Conservative 1: Mismatches Indels 0 111 258 Gaps 207 60 0

Search completed: October 13, 2005, 03:06:57 Job time : 119.425 secs

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Result
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Maximum Match 100%
Listing first 45 summaries
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seq length: 2000000000
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1: /cgn2 6/ptodata/2/pubpaa/PCT NEW PUB.pep:*

2: /cgn2-6/ptodata/2/pubpaa/PCT NEW PUB.pep:*

3: /cgn2-6/ptodata/2/pubpaa/US06 NEW PUB.pep:*

4: /cgn2-6/ptodata/2/pubpaa/US06 NEW PUB.pep:*

5: /cgn2-6/ptodata/2/pubpaa/US07 NEW PUB.pep:*

6: /cgn2-6/ptodata/2/pubpaa/US08 PUBCOMB.pep:*

6: /cgn2-6/ptodata/2/pubpaa/US08 PUBCOMB.pep:*

7: /cgn2-6/ptodata/2/pubpaa/US08 PUBCOMB.pep:*

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11: /cgn2-6/ptodata/2/pubpaa/US09C PUBCOMB.pep:*
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Match
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/ Cgn2 6/ptodata/2/pubpaa/US09B PUBCOMB.pep: */
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US-10-269-805-21

US-10-235-175-80

US-10-371-942-18

US-10-726-332-23

US-10-726-332-23

US-10-726-332-23

US-10-292-088-115

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US-10-120-377-76

US-10-92-196-76
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Sequence 7, Appli
Sequence 21, Appl
Sequence 80, Appl
Sequence 18, Appl
Sequence 19, Appl
Sequence 19, App
Sequence 115, Appl
Sequence 76, Appl
Sequence 76, Appl
Sequence 76, Appl
Sequence 76, Appl
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71,	e 10	؈	10	Sequence 18, Appl	e 51	e 51	e 54	e 35	15	151	136,	12	Sequence 1627, Ap	16	e 13	59	ω •	18	36,	36,	142	Sequence 89, Appl	e 58,	e 78,	e 78,	78,	e 19,		е 59,	110	Sequence 92, Appl	57,	Sequence 63, Appl

## ALIGNMENTS

RESULT 1 US-10-010-729-7

GENERAL INFORMATION:

Sequence 7, Application US/10010729 Publication No. US20030185827A1

APPLICANT: Rodriguez, Moses
APPLICANT: Miller, David J.

APPLICANT: Pease, Larry R.

ITITLE OF INVENTION: Human IgM Antibodies and Diagnostic and
ITITLE OF INVENTION: Therapeutic Uses Thereof Particularly in the Central Nervous
ITITLE OF INVENTION: System
FILE REFERENCE: 1199-1-005CIP2
CURRENT APPLICATION NUMBER: US/10/010,729
CURRENT APPLICATION NUMBER: 09/730,473
PRIOR APPLICATION NUMBER: 09/730,473
PRIOR FILING DATE: 2000-12-05
PRIOR FILING DATE: 2000-15-30
PRIOR FILING DATE: 1990-05-30
PRIOR FILING DATE: 1999-05-28
PRIOR PILING DATE: 1999-05-28
PRIOR PILING DATE: 1997-01-07
PRIOR APPLICATION NUMBER: 08/692,084
PRIOR PILING DATE: 1997-01-07
PRIOR APPLICATION NUMBER: 08/692,084
PRIOR FILING DATE: 1996-08-08
PRIOR FILING DATE: 1996-08-08 NUMBER OF SEQ ID NOS: 80
SOFTMARE: PASTSEQ for Windows Version 4.0
SEQ ID NO 7
LENGTH: 119
TYPE: PAT
ORGANISM: Homo sapiens
US-10-010-729-7 PRIOR APPLICATION NUMBER: 08/236,520 PRIOR FILING DATE: 1994-04-29

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CURRENT APPLICATION NUMBER: US/10/235,175
CURRENT FILING DATE: 2002-09-04
PRIOR APPLICATION NUMBER: US/09/315,926
PRIOR FILING DATE: 1999-05-20
PRIOR APPLICATION NUMBER: EP 99201593.3
PRIOR FILING DATE: 1999-05-20
PRIOR APPLICATION NUMBER: EP 98201693.3
PRIOR APPLICATION NUMBER: EP 98201693.3
PRIOR APPLICATION NUMBER: EP 98201693.3
PRIOR PILING DATE: 1998-05-20
NUMBER: OF SEQ ID NOS: 81
SOFTWARE: Patentin version 3.0
SEQ ID NO 80
LENGTH: 248
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TYPE: PRT
ORGANISM: Homo sapiens
US-10-269-805-21
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US-10-269-805-21
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SEQ ID NO 21
LENGTH: 123
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Publication No. US20030124129A1
GENERAL INFORMATION:
APPLICANT: OLINER, JONATHAN D.
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Best Local :
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Publication No.
                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
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TITLE OF INVENTION: ANGIOPOIETIN-2 SPECIFIC BINDING AGENTS
FILE REFERENCE: A-722
CURRENT APPLICATION NUMBER: US/10/269,805
CURRENT FILING DATE: 2002-10-10
                                                                                                                                                                                                                                                                                        APPLICANT: Es van, Helmuth
APPLICANT: Havenga, Menzo
APPLICANT: Verlinden, Stefan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: US 60/328,604
PRIOR FILING DATE: 2001-10-11
NUMBER OF SEQ ID NOS: 76
                                                                                                                                                                                                                                 TITLE OF INVENTION: TARGETED DELIVERY THROUGH A CATIONIC AMINO ACID TITLE OF INVENTION: TRANSPORTER FILE REFERENCE: 2183-4080US
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Pred. No. 9.6e-50;
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Publication No. US20030223994A1
GENERAL INFORMATION:
APPLICANT: Hoogenboom, Henricus Renerus Jacobus Mattheus
APPLICANT: Reiter, Yoram
TITLE OF INVENTION: MHC-PEPTIDE COMPLEX BINDING LIGANDS
FILE REFERENCE: 10280-034001
                                                                                                                                                                                  Sequence 23, Application US/10726332
Publication No. US20050058649A1
GENERAL INFORMATION:
APPLICANT: Gregory M. Landes
APPLICANT: Mary Haak-Frendscho
APPLICANT: Mary Haak-Frendscho
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Best Local Similarity 87.5%;
Matches 105; Conservative
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Best Local Similarity
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LENGTH: 120
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                                                                                                               APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT:
TITLE OF IN
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CURRENT FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: US 60/358,994
PRIOR FILING DATE: 2002-02-20
NUMBER OF SEQ ID NOS: 121
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
NAME/KEY: misc_feature
OTHER INFORMATION: Description of Artificial Sequence: phage
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                                              OF INVENTION:
Meling
I Xiao Feng
I: Xiao-Chi Jia
IT: Mark R. Nocerini
INVENTION: ANTIBODIES DIRECTED TO PH
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                                                                                                                                                Mary Haak-Frendscho
Ling Chen
Yen-Wah R. Lee
                                                                                                                                                                                                                                                                                                                                                                                 ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAKGVTGS-PTLDYWGQGTLVTVSS
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                                                                                                                                 Meina Liang
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Pred. No. 5.6e-43;
5; Mismatches 9
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Pred. No. 4.1e-43;
2; Mismatches 11; Indels 1;
                                                                  PHOSPHOLIPASE A2
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SOPTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 190
LENGTH: 121
Type.
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PRIOR FILING DATE:
NUMBER OF SEQ ID NOS: 222
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 23
                                                                                                                                               Query Match 87.8%; Score 540; DB 17; Best Local Similarity 87.6%; Pred. No. 4.6e-43; Matches 106; Conservative 2; Mismatches 11;
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Best Local Similarity
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APPLICANT:
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APPLICANT:
APPLICANT:
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CURRENT APPLICATION NUMBER: U2/10/727,155
CURRENT FILING DATE: 2003-12-02
PRIOR APPLICATION NUMBER: 60/430729
PRIOR FILING DATE: 2002-12-02
                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Raffaella Faggioni
APPLICANT: Giorgio Senaldi
APPLICANT: Qiaojuan Jane Su
TITLE OF INVENTION: ANTIHODIES DIRECTED TO TUMOR NECROSIS
TITLE OF INVENTION: FACTOR AND USES THEREOF
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APPLICANT: Jaspal S. Kang
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CURRENT APPLICATION NUMBER: US/10/726,332
CURRENT FILING DATE: 2003-12-02
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TYPE: PRT
ORGANISM: Homo sapiens
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                                                                    1 QVQLVESGGGVVQPGRSLRLSCAASGFTFSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY
ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCAREGDYGGNPYFDYWGQGTLVTVS 120
                       ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAK--GVTGSFTLDYWGQGTLVTVS 118
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Xiao Feng
Scott Klakamp
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Meina Liang
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Palaniswami Rathanaswami
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Pred. No. 4.1e-43;
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Publication No. US20030176674A1
GENERAL INFORMATION:
APPLICANT: Rosen, Craig, et al.
TITLE OF INVENTION: Vascular Endothelial Growth Factor-2
FILE REFERENCE: PF112P8
FILE REFERENCE: PF112P8
CURRENT APPLICATION NUMBER: US/10/120,377
CURRENT FILING DATE: 2002-04-12
FRIOR APPLICATION NUMBER: 60/283,391
FRIOR FILING DATE: 2001-04-13
PRIOR FILING DATE: 2001-09-07
NUMBER: OF SEQ ID NOS: 79
SOFTWARE: Patentin version 3.1
SEQ ID NO 76
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FILE REFERENCE: ABX-PF/3 US
CURRENT APPLICATION NUMBER: US/10/292,088
CURRENT FILING DATE: 2003-03-14
FRIOR REPLICATION NUMBER: 60/348,980
FRIOR FILING DATE: 2001-11-09
NUMBER OF SEQ ID NOS: 147
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 115
LENGTH: 123
TYPE:-PRT
ORGANISH: Homo sapiens
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                                                                                  ; ORGANISM: Homo sapiens US-10-120-377-76
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Matches 105; Conservative
                       Query Match 87.6%;
Best Local Similarity 88.2%;
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Best Local Similarity
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APPLICANT: GLADUB, RONALD
APPLICANT: CORVALAN, JOSE
APPLICANT: JIA, XIAO-CHI
APPLICANT: FENG, XIAO
                                                                                                                     LENGTH: 119
TYPE: PRT
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Score 539; DB 14;
Pred. No. 5.6e-43;
1; Mismatches 13;
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Pred. No. 4.7e-43;
                                       Length 119;
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    Indels
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1 QVQLVESGGGVVQPGRSLRLSCAASGFTFSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY 60

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US-10-980-815-76
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                                                                                 US-10-992-196-76
                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: 60/523,661
PRIOR FILING DATE: 2003-11-21
PRIOR APPLICATION NUMBER: 60/283,391
PRIOR FILING DATE: 2001-04-13
PRIOR APPLICATION NUMBER: 60/317,600
PRIOR FILING DATE: 2001-09-07
PRIOR APPLICATION NUMBER: 10/120,377
PRIOR APPLICATION NUMBER: 10/120,377
PRIOR FILING DATE: 2002-04-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 76, Application US/10992196 Publication No. US20050192429A1 GENERAL INFORMATION:
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Publication No. US20050059117A1
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                     Query Match
Best Local Similarity
                                                                                                                                                                   NUMBER OF SEQ ID NOS:
SEQ ID NO 76
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PRIOR FILING DATE: 2002-04-12
PRIOR APPLICATION NUMBER: 60/283,391
PRIOR FILING DATE: 2001-04-13
PRIOR APPLICATION NUMBER: 60/317,600
PRIOR FILING DATE: 2001-09-07
NUMBER OF SEQ ID NOS: 79
  Matches 105;
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-10-980-815-76
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TITLE OF INVENTION: Vascular Endothelial Growth Factor-2
FILE REFERENCE: PF112P8P1
CURRENT APPLICATION NUMBER: US/10/992,196
CURRENT FILING DATE: 2004-11-19
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CURRENT FILING DATE: 2004-11-04
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TITLE OF INVENTION: Vascular Endothelial Growth Factor-2
FILE REFERENCE: PF112P8
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                                                                                                    ORGANISM: Homo sapiens
                                                                                                                                             LENGTH: 119
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Local Similarity 88.2%;
les 105; Conservative
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  Conservative
                     87.6%;
88.2%;
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Score 539; DB 18; Length 119; Pred. No. 5.6e-43; 1; Mismatches 13; Indels
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Pred. No. 5.6e-43;
1; Mismatches 13
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APPLICANT: BOYLE, WILLIAM
APPLICANT: SULLYAN, John
FILLE OF INVENTION: SELECTIVE BINDING AGENTS OF
FILE REFERENCE: A-633A
CURRENT APPLICATION NUMBER: US/09/791,153A
CURRENT FILING DATE: 2001-07-17
PRIOR APPLICATION NUMBER: 09/511,139
PRIOR APPLICATION NUMBER: 09/511,139
PRIOR FILING DATE: 2000-02-23
NUMBER OF SEQ ID NOS: 154
SOFTWARE: Patentin version 3.0
SEQ ID NO 63
TENTATION 153
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US-10-935-290-100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQ ID NOS: 234
SEQ ID NO 100
LENGTH: 241
TYPE: PRT
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                         Matches 106;
                                                               Query Match
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CURRENT FILING DATE: 2004-09-08
FRIOR APPLICATION NUMBER: PCT/US03/09625
PRIOR FILING DATE: 2003-03-28
PRIOR APPLICATION NUMBER: 60/368,813
PRIOR FILING DATE: 2002-04-01
PRIOR FILING DATE: 2002-04-01
                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Deshpande, Rajendra
APPLICANT: Hitz, Anna
APPLICANT: Boyle, William
APPLICANT: Sullivan, John
                                                                                                                         LENGTH: 113
TYPE: PRT
ORGANISM: Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Baker et al.
TITLE OF INVENTION: Antibodies that Specifically Bind
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILE REFERENCE: PF584P1
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OTHER INFORMATION: scFv protein GMBC683
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Local Similarity 89.1%; hes 106; Conservative
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88.2%;
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                                            Score 538; DB 10;
Pred. No. 6.6e-43;
                         0; Mismatches
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Pred. No. 1.3e-42;
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RESULT 14
US-10-727-155-92
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US-11-021-715-57
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CURRENT APPLICATION NUMBER: US/11/021,715
CURRENT FILING DATE: 2004-12-23
PRIOR APPLICATION NUMBER: PCT/US03/21304
PRIOR FILING DATE: 2003-07-03
PRIOR APPLICATION NUMBER: 60/394,352
PRIOR FILING DATE: 2002-07-03
PRIOR FILING DATE: 2002-09-18
PRIOR FILING DATE: 2002-09-18
PRIOR FILING DATE: 2002-09-18
PRIOR FILING DATE: 2002-09-18
NUMBER OF SEQ ID NOS: 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 92, Application US/107 Publication No. US20050049402A1 GENERAL INFORMATION:
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SEQ ID NO 57
LENGTH: 117
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Best Local :
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APPLICANT:
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             APPLICANT: Raffaella Faggioni
APPLICANT: Giorgio Senaldi
APPLICANT: Qiorjuan Jane Su
TITLE OF INVENTION: ANTIBODIES DIRECTED TO TUMOR NECROSIS
TITLE OF INVENTION: FACTOR AND USES THEREOF
FILE REFERENCE: ADGENIX.073A
CURRENT APPLICATION NUMBER: US/10/727,155
CURRENT FILLING DATE: 2003-12-02
                                                                                                                           APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: John S. APPLICANT: Jaspal APPLICANT: Orit |
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TITLE OF INVENTION: COMPOSITIONS, METHODS AND KITS
TITLE OF INVENTION: AUTOANTIBODIES AND INHIBITORS
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APPLICATION NUMBER: 60/430729
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                                                                                                                                                                                                                                 Craig Pigott
Meina Liang
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Xiao Feng
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Palaniswami Rathanaswami
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    Babcook

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Search completed: October 13, 2005, 03:02:47 Job time : 116.425 secs

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APPLICANT: KELLERMANN, SIRID-AIMEE
APPLICANT: LOW, JOSEBH EDWIN
APPLICANT: MOBLEY, JAMES LESLIE
TITLE OF INVENTION: ANTIBODIES TO M-CSF
FILE REFERENCE: ABS-PF4
CURRENT APPLICATION NUMBER: US/10/938,353
CURRENT APPLICATION NUMBER: 00/502,163
PRIOR APPLICATION NUMBER: 60/502,163
PRIOR FILING DATE: 2003-09-10
NUMBER OF SEQ ID NOS: 117
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-938-353-110
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; Sequence 110, Application US/10938353
; Publication No. US20050059113A1
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NUMBER OF SEQ ID NOS: 320
SOFTWARE: FastSEQ for Windows Version
SEQ ID-NO 92
LENGTH: 121
                                                                                                                                                    Query Match
Best Local Similarity
Matches 105; Conserv
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Best Local Similarity
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APPLICANT: DEVALARAJA, MADHAV NARASIMHA
APPLICANT: FOLTZ, IAN
APPLICANT: HAAK-FRENDSCHO, MARY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Homo
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                        61 ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAKGVTGSPTLDYWGQGTLVTVSS 119
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ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGYS---YFDYWGQGTLVTVSS
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                                                                                                                                                          Conservative
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88.2%; Pred. No. 1.2e-42;
cive 2; Mismatches 9
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Pred. No. 8.8e-43;
3; Mismatches 11
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Result
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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seq length:
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1. /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2. /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3. /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4. /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*

5. /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6. /cgn2_6/ptodata/1/iaa/pcTUS_COMB.pep:*
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Gapop 10.0 , Gapext 0.5
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       GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd
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US-09-315-926A-80
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US-08-759-804A-46
US-09-27-693-46
US-09-472-087-6
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US-09-472-087-6
US-09-456-090A-92
US-09-456-090A-92
US-09-453-234-60
US-09-453-234-108
US-09-472-087-6
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US-09-453-234-106
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US-09-456-090A-106
US-09-456-090A-106
US-09-456-090A-106
US-09-456-090A-106
US-09-456-090A-106
US-09-456-090A-106
US-09-456-090A-68
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46, Appl
46, Appl
46, Appl
163, Appl
63, Appl
64, Appl
64, Appl
66, Appl
60, Appl
60
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81.6	82.1	82.2	82.3	82.4	82.4	82.4	82.8	82.8	82.8	82.8	82.9	83.1	83.4	83.4	83.4	83.4	
116	119	124	125	225	225	125	117	117	117	117	123	115	225	225	225	225	
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US-08-211-202-141	US-09-648-067A-15	US-09-424-840B-16	US-09-240-274-9	US-09-453-234-94	US-09-456-090A-94	US-09-240-274-24	US-09-490-324-24	US-09-490-153-24	US-09-490-070A-24	US-09-025-769B-24	US-09-560-198A-10	US-09-726-219A-167	US-09-453-234-110	US-09-453-234-68	US-09-453-234-56	US-09-456-090A-110	
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#### ALIGNMENTS

RESULT 1 US-09-315-926A-80

Sequence 80, Application US/09315926A Patent No. 6498027

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APPLICANT: Es van, Helmuth
APPLICANT: Havenga, Menzo
APPLICANT: Havenga, Menzo
APPLICANT: Verlinden, Stefan
TITLE OF INVERTION: TARGETED DELIVERY THROUGH A CATIONIC AMINO ACID TRANSPORTER
FILE REFERENCE: 2183-4080US
CURRENT APPLICATION NUMBER: US/09/315,926A
CURRENT APPLICATION NUMBER: BP 99201593.3
PRIOR APPLICATION NUMBER: EP 99201593.3
PRIOR PILING DATE: 1999-05-20
PRIOR APPLICATION NUMBER: EP 98201693.3
PRIOR FILING DATE: 1998-05-20
NUMBER OF SEQ ID NOS: 81
SOPTWARE: Patentin version 3.0
SEQ ID NO 80
LENGTH: 248
TYPE: PRT
ORGANISM: Artificial Sequence
PRATURE:
NAME/KEY: misc feature
OTHER INFORMATION: Description of Artificial Sequence: phage
NAME/KEY: pppTIDE
OTHER INFORMATION: /note="hCAT1 amino acid sequence"
US-09-315:926A-80
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RESULT 2
US-08-331-398A-46
i-Sequence 46, Application US/08331398A
; Patent No. 5608039
; GENERAL INFORMATION:
; APPLICANT: Pastan, Ira
; APPLICANT: Willingham, Mark
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Best Local Similarity
                                                                                                                                                                                                                                           61 ADSVKGRFTISRDNSKNTLYLOMNSLTAXDTAVYYCAKGVTGSPT-LDYWGQGTLVTVSS
                                                                                                                                                                                                             83 ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARGITVTKSRFDYWGQGTLVTVSS 142
                                                                                                                                                                                                                                                                                                 23 QVQLVQSGGGVVQPGRSLRLSCAASGFTFSSYAMHWVRQAPGKGLEWVAVISYDGSNKYY
                                                                                                                                                                                                                                                                                                                           1 QVQLVESGGGVVQPGRSLRLSCAASGPTFSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY
                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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87.5%;
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Pred. No. 1.9e-44;
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                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 4; Length
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FitzGerald, David

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TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9600
TELEPAX: (415) 543-5043
INFORMATION FOR SEQ ID NO: 46:
SEQUENCE CHARACTERISTICS:
LENGTH: 119 amino acids
                                                                                                         Sequence 4
Patent No.
                                                                                                                                                                                                                                                                                                                                                                  Matches 103; Conservative
                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PILING DATE: 30-SEP-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/
FILING DATE: 12-OCT-1990
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Sing
TITLE OF INVENTION:
TITLE OF INVENTION:
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/767,331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                    APPLICANT: Pastan, APPLICANT: Benhar,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: 1..119
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Hunter, Tom
REGISTRATION NUMBER: 38,498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
                                                                                                                            46,
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                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                             6, Application US/08331397B
5981726
                                                                                                                                                                                                                                         ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAKGVTGSPTLDYWGQGTLVTVSS 119
                                                                                                                                                                                                                                                                                                                  QVQLVESGGGVVQPGRSLRLSCAASGFTFSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY 60
                                                                                                                                                                                                                      ADSVKGRETISRDNSKNTLYLOMNSLRAEDTAVYYCARRSARTYYFDYWGQGTLVTVSS 119
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One Market Plaza, Steuart Street Plaza
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                                                                                                                                                                                                                                                                                                                                                                                   86.6%;
Chimeric and Mutationally Stabilized Tumor-
Specific Antibody Fragments, Fusion Proteins, and Uses
Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "Human fetal immunoglobulin
56P1'CL Variable Heavy chain (V-H)"
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Sequence 46, Application US/08759804A Patent No. 5990296

GENERAL INFORMATION:

APPLICANT: Pastan, Ira
APPLICANT: Willingham, Mark
APPLICANT: FitzGerald, David
APPLICANT: Brinkmann, Ulrich
APPLICANT: Pai, Lee

FitzGerald, David Brinkmann, Ulrich

4

TITLE OF INVENTION: Tumor-STITLE OF INVENTION: Fusion NUMBER OF SEQUENCES: 68

Tumor-Specific Antibody Fragments, Fusion Proteins, and Uses Thereof

CORRESPONDENCE ADDRESS

ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco

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RESULT 4
US-08-759-804A-46
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NAME: Hunter, Tom
REGISTRATION NUMBER: 38,498
REFERENCE/DOCKET NUMBER: 0152
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1415) 543-9600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/596,289
FILING DATE: 12-0CT-1990
ATTORNEY/AGENT INFORMATION:
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MEDIUM TYPE: Floppy disk
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LENGTH: 119 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: One Mainten
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                    LOCATION: 1..119
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US 0 FILING DATE: 30-SEP-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/331,397B FILING DATE: 28-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: 11
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                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY:
                                                                                 51
                                                                                                                        61 ADSVKGRFTISRDNSKNTLYLOMNSLTAXDTAVYYCAKGVTGSPTLDYWGQGTLVTV8S 119
                                                                                                                                                                                                     1 QVQLVESGGGVVQPGRSLRLSCAASGFTFSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  94105-1492
                                                                                 ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCARRSARTYYFDYWGQGTLVTVSS
                                                                                                                                                                QVELVESGGGVVQPGRSLRLSCAASGFTFSSYAMHWVRQAPGKGLEWVAVISYDGSNKYY 60
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                                                                                                                                                                                                                                                86.3%; Score 531; DB 2;
86.6%; Pred. No. 1.1e-43;
Live 3; Mismatches 13
                                                                                                                                                                                                                                                                                                                                                    /note= "Human fetal immunoglobulin
56P1'CL Variable Heavy chain (V-H)"
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk

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94111-3834

California

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                                                                                                                                                                                                                                                                                  Sequence 46,
                                                                                                                                                                                                                                                               Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches 103; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                     GENERAL INFORMATION:

APPLICANT: BENHAR, Ira

APPLICANT: BENHAR, Itai

APPLICANT: PADLAN, Eduardo A.

APPLICANT: JUNG, Sun-Hee

APPLICANT: LEE, Byungkook

TITLE OF INVENTION: HUMANIZED TUMOR-SPECIFIC ANTIBODY

TITLE OF INVENTION: FRAMMENTS, FUSION PROTEINS, AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 03-DEC-1996
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/331,398
FILING DATE: 28-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/767,331
APPLICATION NUMBER: US 07/767,331
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Weber, Ellen L.
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 01:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 30-SEP-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/596,289
FILING DATE: 12-OCT-1990
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/759,804A
                                                           CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: protein
                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: 1..119
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
              STREET:
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                                      ADDRESSEE:
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San Francisco
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                 Steuart Street Tower,
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                 Townsend and Townsend Khourie and Crew suart Street Tower, One Market Plaza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       86.3%;
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56P1'CL Variable Heavy chain (V-H)"
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Pred. No. 1.1e-43;
3; Mismatches 13
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                                                                                 FILE REFERENCE: ABX-PF1
CURRENT APPLICATION NUMBER: US/09/472,087
CURRENT FILING DATE: 1999-12-23
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                                                                                                                                     APPLICANT: HANSON, DOUGLAS C.
APPLICANT: MUBLER, EILLEN E.
APPLICANT: HANKE, JEFFREY H.
APPLICANT: GILMAN, STEVEN C.
APPLICANT: DAVIS, C. GEOFFREY
APPLICANT: DAVIS, C. GEOFFREY
APPLICANT: CORVALAN, JOSE R.
TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES TO CTLA-4
PRIOR APPLICATION NUMBER: 60/113,647
PRIOR FILING DATE: 1998-12-23
NUMBER OF SEQ ID NOS: 147
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (415) 543-5043
INFORMATION FOR SEQ ID NO: 46:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 30-SEP-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 0
FILING DATE: 12-OCT-1990
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PILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/331,396
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 543-9600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: Protein LOCATION: 1..119 OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE:
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56Pl'CL VH region"
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; ORGANISM: Homo sapiens
US-09-472-087-63
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US-09-472-087-63
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; ORGANISM: Homo sapiens
US-09-472-087-1
                                                                                                                                                                                                                                                                  US-09-472-087-64
              GENERAL INFORMATION:
APPLICANT: HANSON, DOUGLAS C.
APPLICANT: MUELLER, EILLEN E.
APPLICANT: MUELLER, EILLEN E.
APPLICANT: HANKE, JEFFREY H.
APPLICANT: GLIMAN, STEVEN C.
APPLICANT: DAVIS, C. GEOFFREY
APPLICANT: DORVALAN, JOSE R.
TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES TO CTLA-4
FILE REFERENCE: ABX-PF1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 1
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Best Local Similarity 86.6%;
Matches 103; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                        Sequence 64, Application US/09472087 Patent No. 6682736
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CURRENT FILING DATE: 1999-12-23
PRIOR APPLICATION NUMBER: 60/113,647
PRIOR FILING DATE: 1998-12-23
NUMBER OF SEQ ID NOS: 147
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: HANSON, DOUGLAS C.
APPLICANT: MUELLER, BILLEN E.
APPLICANT: HANKE, JEFFREY H.
APPLICANT: GILMAN, STEVEN C.
APPLICANT: GILMAN, STEVEN C.
APPLICANT: DAVIS, C. GEOFREY
APPLICANT: CORVALAN, JOSE R.
TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES TO CTLA-4
FILE REFERENCE: ABX-PF1
CURRENT APPLICATION NUMBER: US/09/472,087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 463
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86.6%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 525.5; DB 4
Pred. No. 1.6e-42;
2; Mismatches 13
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Pred. No. 1.6e-
2; Mismatches
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US-09-472-087-64
                                                                                                                                        ; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-07-942-245-35
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PRIOR APPLICATION NUMBER: 60/113,647
PRIOR FILING DATE: 1998-12-23
NUMBER OF SEQ ID NOS: 147
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 85.4%;
Best Local Similarity 86.6%;
Matches 103; Conservative
                                                                      Matches
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                                                                                    Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                             TELEPHONE: (202) 293-70
TELEFAX: (202) 293-7860
TELEX: 6491103
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 20037-3202
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: HP 9000/700 Workstation
                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/07/942,245
FILING DATE: 09-SEP-1992
CLASSIFICATION: 530
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 293-7060
                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: UNIX SOFTWARE: In house CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: GUILD,
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 120 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 522
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sughtue, Mion, Zinn, Macpeak & Seas
STREET: 2100 Pensylvania Avenue, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Washington STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 QVQLVESGGGVVQPGRSLRLSCVASGFTFSSHGMHWVRQAPGKGLEWVAVIWYDGRNKYY 79
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                    QVQLVESGGGVVQPGRSLRLSCAASGFTFSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY 60
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QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYAMHWVRQAPGKGLEWVAVISYDGSNKYY
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                                                                                    85.4%;
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Michael A.
Braydon C.
SURFACE RESIDUE VENEERING OF RODENT
ANTIBODIES
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Stephen M.J
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Pred. No. 1.6e-42;
                                                                    Score 525; DB 1;
Pred. No. 4.2e-43;
1; Mismatches 13
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GENERAL INFORMATION:

APPLICANT: Buechler, Joe

APPLICANT: Valkirs, Gunars

APPLICANT: Valkirs, Gunars

APPLICANT: Lonberg, Nils

FITTLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS

FILE REFERENCE: 020015-000200US

CURRENT APPLICATION NUMBER: US/09/456,090A

CURRENT FILING DATE: 1999-12-06

NUMBER OF SEQ ID NOS: 110

SOFTWARE: PATEENTIN Ver. 2.1

SEQ ID NO 92

LENGTH: 225

TYPE: PRT

ORGANISM: Homo sapiens

OTHER INFORMATION: M2-11H
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US-09-456-090A-92
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US-09-456-090A-60
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Best Local Similarity 85.7
Matches 102; Conservative
                                                                                                                                                                      Query Match 84.7%;
Best Local Similarity 85.7%;
Matches 102; Conservative
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LENGTH: 225
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APPLICANT: Lonberg, WAIN
TITLE OF INVENTION: HUMAN ANTIBODIES AS DIAGNOSTIC REAGENTS
FILE REFERENCE: 020015-000200US
CURRENT APPLICATION NUMBER: US/09/456,090A
CURRENT FILING DATE: 1999-12-06
NUMBER OF SEQ ID NOS: 110
SOFTWARE: Patentin Ver. 2.1
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APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff
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ORGANISM: Homo sapiens
OTHER INFORMATION: M1-5H
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6680209
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                          ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAKGVTGSPTLDYWGQGTLVTVSS 119
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ADSVKGRFTISRDNSKNTLYLOMNSLRAEDTAVYYCAR -- DGIGYFDYWGQGTLVTVSS 117
                                                                                  QVQLVESGGGVVQPGRSLRLSCAASGFTFSYYGMHWVRQAPGKGLEWVTLITYDGDNKYY 60
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85.7%; Pred. No. 2e-42;
                                                                                                                                                                      2; Mismatches
                                                                                                                                                                                          Score 521; DB 4; Length 225; Pred. No. 2e-42;
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US-09-453-234-60
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    Matches 102;
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Matches 102; Conservative
                                            Query Match
                                                                                                                                                                                            SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 92
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CURRENT FILING DATE: 1999-12-01
                                                                                                                                                                                                                                                        TITLE OF INVENTION: Human Antibodies FILE REFERENCE: 020015-000110US CURRENT APPLICATION NUMBER: US/09/453,234 CURRENT FILING DATE: 1999-12-01 PRIOR APPLICATION NUMBER: US 60/157,415 PRIOR FILING DATE: 1999-10-02
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APPLICANT: Biosite Diagnostics, Inc.
APPLICANT: GenPharm International
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APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff
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APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff
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APPLICANT: Biosite Diagnostics, Inc.
APPLICANT: GenPharm International
TITLE OF INVENTION: Human Antibodies
                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 112
                                                                                                      TYPE: PRT
ORGANISM: Homo sapiens
OTHER INFORMATION: M2-11H
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 225
TYPE: PRT
ORGANISM: Homo sapiens
OTHER INFORMATION: M1-5H
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                                                                                                                                                                           LENGTH: 225
                        Match 84.7%;
Local Similarity 85.7%;
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    Conservative
2;
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Pred. No. 2e-42;
2; Mismatches
                      Score 521; DB 4; Length 225; Pred. No. 2e-42;
    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 4; Length 225;
    13; Indels
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  Gaps
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1 QVQLVESGGGVVQPGRSLRLSCAASGFTFSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY 60

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APPLICANT: Buechler, Joe
APPLICANT: Walkirs, Gunars
APPLICANT: Gray, Jeff
APPLICANT: Lonberg, Nils
FILE REFERENCE: 020015-000220US
FILE REFERENCE: 020015-000220US
CURRENT APPLICATION NUMBER: US/09/456,090A
CURRENT FILING DATE: 1999-12-06
NUMBER OF SEQ ID NOS: 110
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 108
LENGTH: 225
TYPE: PAT
ORGANISM: Homo sapiens
OTHER INFORMATION: M2-34H
US-09-456-090A-108
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US-09-453-234-108
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                                                                                                                              ; TYPE: PRT
; ORGANISM: Homo sapiens
; OTHER INFORMATION: M2-34H
US-09-453-234-108
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APPLICANT: Valkirs, Gunars
APPLICANT: Gray, Jeff
APPLICANT: Lonberg, Nils
APPLICANT: Lonberg, Nils
APPLICANT: Biosite Diagnostics, Inc.
APPLICANT: GenPharm International
TITLE OF INVENTION: Human Antibodies
FILE REFERENCE: 020015-000110US
CURRENT APPLICATION NUMBER: US/09/453,234
CURRENT FILING DATE: 1999.12-01
                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: US 60/157,415
PRIOR FILING DATE: 1999-10-02
NUMBER OF SEQ ID NOS: 112
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 108
LENGTH: 225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 108, Application US/09453234
Patent No. 6794132
GENERAL INFORMATION:
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Best Local Similarity 85.7
Matches 102; Conservative
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                                                              Query Match
Best Local Similarity
                                          Matches 102; Conservative
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1 QVQLVESGGGVVQPGRSLRLSCAASGFTFSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY 60
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                                                                84.6%; Score 520; DB 4; Length 225; 85.7%; Pred. No. 2.5e-42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       84.6%; Score 520; DB 4; Length 225; 85.7%; Pred. No. 2.5e-42;
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                                          2; Mismatches
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              61 ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAKGVTGSPTLDYMGQGTLVTVSS 119
                                                 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDWIG--YFDYWGQGTLVTVSS 117
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Search completed: October 13, Job time: 20.4077 secs 2005, 02:45:19

#### **Protein Sequence Searches - February 2005**

All of the sequence databases on ABSS have recently been updated.

- Please note that the curators of the UniProt database have purged some temporary accession numbers from the most recent version of UniProt. These sequences have been assigned new permanent accession numbers. The new UniProt record may not contain the previous temporary accession number.
- If you encounter an accession number from an older search run against UniProt (results file extension .rup) that can no longer be found in the database, the permanent record with the new accession number can be found by searching the old accession number in the UniProt Protein. Archive database (UniPARC) at:

http://www.pir.uniprot.org/database/archive.shtml

If you have any questions regarding this information or your results, please contact any STIC searcher.

When submitting sequence search results for scanning into IFW, please include a copy of this attachment to assist any future Examiners or members of the public who may encounter UniProt temporary accession numbers.

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Total number of hits satisfying chosen parameters:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 QVQLVESGGGVVQPGRSLRL.....VTGSPTLDYWGQGTLVTVSS 119
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                Q6mzv6
Q6mzv7
Q9hcc1
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Q9u191
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P01772
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Q8wu38
P01781
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  sapien
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Q8WUK1
ID Q8WU
     RX MEDLINE-2238857; PubMed=12477932; DOI=10.1073/pnas.242603899;
RA Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschall S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschanko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., WcKernan K.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Raha S.S., Morley K.C., Hale S., Garcia A.M., Gybur S.A.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Pahey J., Helton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Pahey J., Halton E., Ketteman M., Madan A., Rodrigues S., Sanchez A.,
RA Hilaseley R.W., Touchman J.W., Green E.D., Dickson M.C.,
RA Kraywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Tones S. T. Marya M.A.
Strausberg R.;
Strausberg R.;
Submitted (DEC-2001) to the EN Submitted (DEC-2001) to the EN EMBL; BCC02040. AAH20240.1; -.
PIR, F36005; F36005.
PIR, PH6405; PH642.
PIR, PH1643; PH1643.
PIR, PH16445; PH16445.
PIR, PH1645; PH1646.
PIR, S11116.
PIR, S31116; S31116.
PIR, S311116; S311116.
PIR, S311119; S311119.
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QBWUK1;
QBWUK1;
01-MAR-2002 (TrEMBLrel. 20, C
01-MAR-2002 (TrEMBLrel. 20, L
01-MAR-2004 (TrEMBLrel. 26, L
                                                                                                                                                                                                                                               Jones S.J., Marra M.A.; "Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences."; Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
TISSUE=Primary B-Cells;
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Homo sapiens (Human).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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TISSUE=Primary B-Cells;
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HV3B HMAN
HV16 MOUSE
Q8N5K4
Q6N6C4
Q6P6C4
Q99X6A4
Q6N034
Q91Z05
Q92C07
Q6PDB8
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Last annotation update)
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Q6p6c4
Q99ka4
Q99ka4
Q91z05
Q91z05
Q920e7
Q920e7
Q6pdb8
Q6mzx9
Q96k68
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Result No.

Score

Match

Length

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Q9UL84 HV3G HUMAN Q9Y509 HV3J HUMAN Q9UL71

Q9UL90 Q65ZC9

Q96BB9 HV3H H Q6PI81

HUMAN

Q6GMX2

Q6PJ95 HV3I\_HUMAN Q6PJĀ4

Q6MZV6 Q6MZV7 Q9HCC1 Q6GMY2

Q6N092 HV3L\_HUMAN

Q8NCI6 Q9UL91 Q6N089 HV3K HUMAN Q6IN78

Q8WU38 HV3T\_HUMAN

525 525 526 513 490.5 482.5 482.5 482.5 482.5 482.5 464.5 464.5 464.5 464.5 464.5 464.5 464.5 464.5 464.5 464.5 465.5 46

887. 805. 807.

Q9UL72 Q6MZU6

Database

UniProt\_03:\*
1: uniprot\_sprot:\*
2: uniprot\_trembl:\*

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Minimum DB Maximum DB

seq length: 0
seq length: 2000000000

Scoring table: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5

Perfect score:

US-10-010-729A-7 615

OM protein -

protein search, using sw model

October 13, 2005, 02:45:26;

Searched:

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Q9UL90 I

Q9UL90 I

AC Q9UL90;

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DT 01-MAY-2000

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Q9UL93;
01-MAY-2000 (TrEMBLrel. 1
01-MAY-2000 (TrEMBLrel. 1
01-MAY-2004 (TrEMBLrel. 2
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SEQUENCE
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SEQUENCE FROM N.A.
MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531;
Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     HSSP; P01861; IADQ.
Pfam; PF07654; C1-set; 4.
SMART; SM00406; IGv; 1.
PROSITE; PS50835; IG_LIKE; 5.
PROSITE; PS00290; IG_MHC; UNKNOWN
SEQUENCE 613 AA; 67295 MW; 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HSSP; P01772; 2FB4.
InterPro; IPR007710; Ig-like.
InterPro; IPR007936; Ig-v.
SMART; SM00406; IGv; 1.
PROSITE; P850835; IG_LIKE; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Clin. Immunol. Immunopathol. 87:184-192(1998)
EMBL; AF035021; AAD56257.1; -.
PIR; PH1644; PH1644.
PIR; PL0120; PL0120.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Bulheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAR-2004 (TrEMBLrel. 26, Last annotation update)
Myosin-reactive immunoglobulin heavy chain variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Myosin-reactive autoantibodies
                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
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116 AA;
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RESULT 4
Q65ZC9
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Best Local S
Matches 103
Query Match
Best Local Sim
Matches 101;
                                                                                                                                       Pfam; PF00047; 19; 2. T
SMART; SM00406; IG; 2.
SMART; SM00406; IGv; 2.
PROSITE; PS50835; IG_LIKE; 2
NON_TER 1 1
NON_TER 240 240
SEQUENCE 240 AA; 25569 MP
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InterPro; IPR003596; Ig_v.
SMART; SM00406; IGv; 1.
PROSITE; P850835; IG_LIKE; 1.
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SEQUENCE FROM N.A.
MEDLINE=9877139; PubMed=9614934; DOI=10.1006/clin.1998.
Mu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Kontermann R.E., Wing M.G., Winter G.; "Complement recruitment using bispecific Nat. Biotechnol. 15:629-631(1997).
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-OCT-2004 (TrEMBLrel. 28, 25-OCT-2004 (TrEMBLrel. 28, 25-OCT-2004 (TrEMBLrel. 28,
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                            EMBL; Y13056; CAA73499.1; -.
InterPro; IPR003599; Ig.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A. STRAIN=Clq/7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Single-chain
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PIR; S78486; S78486.
HSSP; P01772; 2FB4.
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Young D.C.;
"Myosin-reactive autoantibodies in rheumatic carditis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Myosin-reactive immunoglobulin heavy chain variable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRAIN=Clq/7;
MEDLINE=97362799;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Name=scFv;
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03; Conservative
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Wing M.G., Winter
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Last
       Score 513; DB Pred. No. 2.3e 5; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Created)
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Pred. No. 8.1e-49;
2; Mismatches 8
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Best Local S
Matches 98
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Q9UL84;
01-MAY-2000
                                                                                                                             21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
05-JUL-2094 (Rel. 44, Last annotation updat
Ig heavy chain V-III region CAM.
Homo sapiens (Human)
                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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InterPro; IPR003596; Ig-v.
SMART; SM00406; IGv; 1.
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EMBL; AF035030; AAD56266.1; -.
HSSP; P01772; 2FB4.
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MEDLINE=98277139; PubMed=9614934; DOI=10.1006/clin.1998.4531;

Wu X., Liu B., Van der Merwe P.L., Kalis N.N., Berney S.M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Ol-WAY-2000 (TrEMBLrel. 13, Created)
Ol-MAY-2000 (TrEMBLrel. 13, Last sequ
Ol-CCT-2000 (TrEMBLrel. 25, Last and
Myosin-reactive immunoglobulin heavy
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Mammalia; Eutheria;
WEDLINE=81013859;
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                        EQUENCE.
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122 AA;
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                                                                                                                                                                                                                                                                              STANDARD;
  PubMed=6774332
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13579 MW;
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Primates;
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Last annotation update)
lin heavy chain variable region
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GO; GO:0003923; F:antigen binding; NAS.
GO; GO:0006955; P:immune response; NAS.
InterPro; IPR007110; Ig-like.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
Pfam; PP00047; Ig; 1.
SMART; SM00406; IGv; 1.
ENOSITE; PS50835; IG_IKE; 1.
Direct protein sequencing; Immunoglobulin
Pyrrolidone carboxylic acid.
Pyrrolidone carboxylic acid.
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Ig-like.
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GO; GO:0016066; P:cellular defer
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PROSITE; PS50835; IG_LIKE; 1.
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Leukemia 9:1948-1953(1995).
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MEDLINE=96071149, PubMed=7475288;
MEDLINE=96071149, PubMed=7475288;
Cao J., Vescio R.A., Rettig M.B., Hor
Lichtenstein A.K., Berenson J.R.;
"A CD10-positive subset of malignant."
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HSSP; P01842; 1AQK.
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Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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HSSP; P01772; 2FB4.
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-i- SIMILARITY: Contains 1 immunoglobulin-like domain
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21-JUL-1986 (Rel. 01, Last sequence up
05-JUL-2004 (Rel. 44, Last annotation
Ig heavy chain V-III region HIL.
Homo sapiens (Human).
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Biochemistry 18:553-560(1979).
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MEDLINE=79124695;
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SMART; SM00406; IGv; 1.

PROSITE; PS50835; IG_LIKE; 1.

Direct protein sequencing; Immunoglobulin V

Pyrrolidone carboxylic acid.

DOMAIN 1 112 Ig-like.
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"Amino acid sequence of the VH region
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30; G0:0003823; F:antigen binding; NAS.
30; G0:0006955; P:inmune response; NAS.
InterPro; IPR007110; Ig-like.
InterPro; IPR003596; Ig_v.
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                                                                                              GTLVTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GTLVTVSS
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                                                      ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAK--GVTGSPTLDYWGQGTLVTVS
                                         QVKLVQAGGGVVQPGRSLRLSCIASGFTFSNYGMHWVRQAPGKGLEWVAVIWYNGSRTYY
                                                                                                                                                                      121
121 AA;
121
                     119
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                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                           PubMed=420800;
ez de Castro J.A.,
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                                                                                                                                                                     121
13566 MW;
                                                                                                                                                                                                                                                                                                                                                                  This chain was isolated from an IgG1 myeloma
                                                                                                                                       78.4%;
76.0%;
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                                                                                                                                                                                                                                                                                                                                             immunoglobulin-like
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Pred. No. 2.7e-
6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6
                                                                                                                                        Score 482;
Pred. No.
                                                                                                                                                                                          Ig-like.
Pyrrolidone carboxylic acid.
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                                                                                                                                                                        480FC53610EF5DAB CRC64;
                                                                                                                              Mismatches
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on of human myeloma
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                                                                                                                                        2.5e-44;
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                                                                                                                              16;
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  Q6GMX2
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Q6GMX2;
05-JUL-2004 (TrEMBLrel. 2
05-JUL-2004 (TrEMBLrel. 2
05-JUL-2004 (TrEMBLrel. 2
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Q9UL71;
Q1-MAY-2000
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-OCT-2003 (TrEMBLrel. 25, Last annotation updat
Myosin-reactive immunoglobulin heavy chain variab
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens (Human)
Eukaryota; Metazoa;
Mammalia; Eutheria;
Strausberg R.L., Feingold B.A., Grouse L.H., Derge J.G.,
Strausberg R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
Rlausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
Rltschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Heleh F.,
Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.,
Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
                                                                                                                                                                                                                                                                                                                                                                                  Hypothetical protein. Homo sapiens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wu X., Liu B., Van Young D.C.,
                                                                                                                                                                                                                                                                                                                                      Bukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
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                                                                                                                                                                                                     TISSUE=59leen;
MEDLINE=22388257; PubMed=12477932; DOI=10.1073/pnas.242603899;
                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                        NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     InterPro;
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Mu X., Liu B., Van der Merwe P.L., Kalis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADSVKGRFTISRDNSKNTLYLOMNSLTAXDTAVYYCAKG--VTGSPTLDYWGQGTLVTVS
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121 AA;
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13154 MW;
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Primates;
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Primates;
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77.7%;
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27,
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Last sequence
Last annotation
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Pred. No. 1.8e-43;
5; Mismatches 20;
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                                                                                                                                                                                                                                                                                                                                        Craniata; Vertebrata; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2F045CCFA5D50736 CRC64;
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annotation update)
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N.N., Berney S.M.,
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Best Local S
Matches 93
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InterPro; IPR007110; Ig-1ike.
InterPro; IPR007110; Ig-c1.
InterPro; IPR003597; Ig_c1.
InterPro; IPR003595; Ig_MHC.
InterPro; IPR003595; Ig_v.
Pfam; PP00647; C1-set; Z.
Pfam; PP00047; Ig; 3.
SMART; SM00409; IG; 4.
SMART; SM00407; IGc1; 2.
SMART; SM00407; IGc1; 1.
PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS50835; IG_LIKE; 4.
PROSITE; PS50835; IG_MHC; UNKNOWN_1.
MEDIJUE-230827; PubMed=12477932; DOI=10.1073/pnas.242603899; Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G., Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D., Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K., Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F., Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L., Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E., Brownstein M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C., Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
                                                                                                                                                                                                                                                                      Q96BB9;
01-DEC-2001 (TrEMBLrel.
01-DEC-2001 (TrEMBLrel.
01-MAR-2004 (TrEMBLrel.
                                                                                                                                               SEQUENCE FROM N.A. TISSUE=Primary B-C
                                                                                                                                                                                       Eukaryota; Metazoa;
Mammalia; Eutheria;
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                         Q96BB9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Submitted (JUN-2004) to the EMBL/GenBank/DDBJ databases. EMBL; BC073771; AAH73771.1; -. InterPro; IPR003599; Ig.
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                                                                                                                                                                                                                                          Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jones S.J., Marra M.A.; "Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.";
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TISSUE=Spleen;
                                                                                                                                                                                                                                                                                                                                                                                                                        140
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADSVKGRFTISRDNAKNTLYLQMNSLRGEDAAVYYCARGFVSLPRSTLDIWGQGTMVTVS
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493 AA; 5
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                                                                                                                                               B-Cells;
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76.9%;
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Primates;
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19,
26,
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Last annotation update)
                                                                                                                                                                                                                                                                                                        Created)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 471; DB 2;
Pred. No. 1.9e-42;
                                                                                                                                                                                                           Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                         PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             55B999305B286203 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2;
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                                                                                                                                                                                                                                                                                                                                                                             RESULT 12
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3H HUMAN
3H HUMAN
STANDARD;
P01769;
P01769;
T 21-JUL-1986 (Rel. 01, Created)
T 21-JUL-1986 (Rel. 01, Last seq
)T 05-JUL-2004 (Rel. 44, Last ann
)T 05-JUL-2004 (Rel. 44, Last ann
)T 05-JUL-2004 (Rel. 44, Last ann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 94
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PIR; S05271; S05271

PIR; S24560; S24260.

H88P; P01861; IADQ.

InterPro; IPR007110; Ig-1ike.
InterPro; IPR003597; Ig_c1.
InterPro; IPR003596; Ig_MCC.
InterPro; IPR003596; Ig_WC.

InterPro; IPR003596; Ig_V.

R InterPro; IPR003596; Ig_V.

R InterPro; IPR003596; Ig_V.

R INTERPRO; IPR003596; Ig_V.

R SMART; SM00406; IGV; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W., Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A., Yallalon D.K., Metheman M., Madan A., Rodrigues S., Sanchez A., Whiting M., Madan A., Young A.C., Shevchenko Y., Bouffard G.G., Blakesley R.W., Touchman J.W., Green B.D., Dickson M.C., Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S., Krzywinski M.I., Skalska U., Smallus D.B., Schnerch A., Schein J.B.,
HSSP; P01772; 2FB4.
GO; GO:0005576; C:extracellular; NAS.
GO; GO:0003823; F:antigen binding; NAS
                                                                                          Plorent G., Lehman D., Putnam F.W.; Plorent G., Lehman D., Putnam F.W.; The switch point in mu heavy chains of hum Biochemistry 13:2482-2498 (1974).

-I- MISCELLANEOUS: This chain was isolated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Jones S.J., Marra M.A.; "Generation and initial analysis of more than 15,000 and mouse CDNA sequences."; Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; I
Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                 Homo sapiens (Human)
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PROSITE; PS00290; IG_MHC; UNKNOWN
SEQUENCE 597 AA; 65039 MW; 4FC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Submitted
                                               PIR; A02052; M3HUGA.
                                                            macroglobulin.
-!- SIMILARITY: Contains 1 immunoglobulin-like
                                                                                                                                                                       SEQUENCE.
                                                                                                                                                                                                  NCBI_TaxID=9606;
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TISSUE=Primary B-Cells;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Bosak S.A., McEwan P.J., McKernan K.J., Malek J.A., Gunaratne P.
Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk
                                                                                                                                                        MEDLINE=74175307; PubMed=4208843;
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                                                                                                                                                                                                                                                                                                                                                                                                                            SSALA
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                                                                                                                                                                                                                                                                01, Last sequence update)
44, Last annotation updat
III region GA.
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75.2%;
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; Pred. No. 5e-42;
10; Mismatches
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4FCA3AD8ECE263D9 CRC64;
                                                                                                                         of human
                                                                                                                                                                                                                                                                                                                                                122
                                                                                                                                                                                                                                                                                  update)
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                                                                                            from a Waldenstrom's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2; Length 597;
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                                                                                                                           IgM immunoglobulins.";
                                                                domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                    Euteleostomi;
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QGPIB 1

ID QGPIB 1

ID QGPIB 2

GGPIB 2

GGPIB 3

CGPIB 3

CGPIB 3

CGPIB 4

CGPIB 4

CGPIB 4

CGPIB 4

CGPIB 6

CGC MAMMMM 6

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RX MEDLINE=22388557; PubMede=12477932; DOI=10.1073/pnas.242603899;

RX MEDLINE=22388557; PubMede=12477932; DOI=10.1073/pnas.242603899;

RX Strausberg R.L., Feingold E.A., Grouse L.H., Derge J.G.,

RX Klausner R.D., Collins P.S., Wagner L., Shenmen C.M., Schuler G.D.,

RX Klausner R.D., Collins P.S., Wagner L., Shenmen C.F., Bhat N.K.,

RX Alteschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RX Alteschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,

RX Alteschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,

RX Alteschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hong L.,

RX Stapleton M., Soares M.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,

RX Stapleton M.J., Usdin T.B., Toshiyuki S., Carninci P., Prange C.J.,

RX Ala S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RX Ala S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,

RX Ala S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.W.,

RX Ala S.S., Loquellano N.A., Sodergren B.J., Lu X., Gibbs R.A.,

RX Alla S., Morley K.C., Hale S., Garcia A.M., Gay L.J., Hullyk S.W.,

RX Alla S.S., Loquellano N.A., Sodergren B.J., Lu X., Gibbs R.A.,

RX Araywinski M., Touchman J.W., Green B.D., Dickson M.C.,

RX RX Araywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.B.,

RX Araywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.B.,

RX Araywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.B.,

RX Araywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.B.,
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Best Local S
Matches 85
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hypothetical protein.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-JUL-2004 (TrEMBLrel.
05-JUL-2004 (TrEMBLrel.
05-JUL-2004 (TrEMBLrel.
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SMART; SM00406; IGv; I.
PROSITE; PS50835; IG; II.
Direct protein sequencing; Immunoglobulin V region;
Pyrrolidone carboxylic acid.
DOMAIN 1 112 Ig-like.
Strausberg R.;
Submitted (DEC-2002) to the EMBL/GenBank/DDBJ databases.
Submit BC041037; AAH41037.1; -.
EMBL; BC041037; AAH41037.1; -.
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InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                      "Generation and initial analysis of more than 15,000 full-length human and mouse cDNA sequences.";
Proc. Natl. Acad. Sci. U.S.A. 99:16899-16903(2002).
                                                                                                                                               SEQUENCE FROM N.A.
TISSUE=Primary B-Cells;
                                                                                                                                                                                                                                                                                                                              Jones S.J., Marra M.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RES
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13166
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27,
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; Pred. No. 9.3e
20; Mismatches
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Last sequence update)
Last annotation update)
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Pyrrolidone carboxylic acid.
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InterPro; IPRO07110; Ig-1ike.

DR InterPro; IPRO07110; Ig-1ike.

DR InterPro; IPRO03597; Ig cl.

DR InterPro; IPRO03596; Ig-MHC.

DR InterPro; IPRO03596; Ig-V.

DR Pfam; PR07654; Cl-set; 3.

DR SMART; SM00409; IG; 2.

DR SMART; SM00407; IGcl; 3.

R SMART; SM00406; IGv; 1.

R PROSITE; PS50835; IG LIKE; 4.

R PROSITE; PS50835; IG MHC; UNKNOWN 2.

# Hypothetical protein.

# SEQUENCE 478 AA; Englished.
Q66795
ID Q66799
AC Q66799
AC Q66799
AC Q66799
AC Q66799
AC Q66799
AC Q66799
O5-JU
DT Q5-JU
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RA Strausberg R.L., Feingold B.A., Grouse L.H., Derge J.G.,
RA Klausner R.D., Collins F.S., Wagner L., Shenmen C.M., Schuler G.D.,
RA Altschul S.F., Zeeberg B., Buetow K.H., Schaefer C.F., Bhat N.K.,
RA Altschul S.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Hopkins R.F., Jordan H., Moore T., Max S.I., Wang J., Hsieh F.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Rubin G.M., Hong L.,
RA Diatchenko L., Marusina K., Farmer A.A., Gunaratne P.H.,
RA Diatchenko L., Moley N.B., Bonaldo M.F., Casavant T.L., Scheetz T.E.,
RA Brownstein M.J., Uedin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Brownstein M.J., Uedin T.B., Toshiyuki S., Carninci P., Prange C.,
RA Raha S.S., Loquellano N.A., Peters G.J., Abramson R.D., Mullahy S.J.,
RA Robesk S.A., McZwan P.J., McKernan K.J., Malek J.A., Gunaratne P.H.,
RA Richards S., Worley K.C., Hale S., Garcia A.M., Gay L.J., Hulyk S.W.,
RA Villalon D.K., Muzny D.M., Sodergren E.J., Lu X., Gibbs R.A.,
RA Pahey J., Helton B., Ketteman M., Wadan A., Rodrigues S., Sanchez A.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Rodriguez A.C., Grimwood J., Schmutz J., Myers R.M., Butterfield Y.S.,
RA Roywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Roywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
RA Roywinski M.I., Skalska U., Smailus D.E., Schnerch A., Schein J.E.,
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Best Local S
Matches 93
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Q6FJ95;
05-JUL-2004 (TrEMBLrel. 2:
05-JUL-2004 (TrEMBLrel. 2:
05-JUL-2004 (TrEMBLrel. 2:
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hypothetical protein. Homo sapiens (Human).
                                                                                               SEQUENCE FROM N.A.
TISSUE=Primary B-Cells;
                                                                                                                                                                                                                           Jones S.J., Marra M.A.; "Generation and initial analysis and mouse cDNA sequences.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9606;
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l Similarity 72.1%;
93; Conservative
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478 AA; 52666 MW; 17BED38D917970D6 CRC64;
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27,
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Last sequence update)
Last annotation updat
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Pred. No. 5e-42;
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2001) to the AAH19046.1;

EMBL/GenBank/DDBJ databases

InterPro; IPR003599; InterPro; IPR007110;

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RESULT 15
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Best Local Similarity
Matches 96; Conserv
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InterPro; IPRO03597; Ig-c1.
InterPro; IPRO03597; Ig-c1.
InterPro; IPRO03596; Ig-MHC.
InterPro; IPRO03596; Ig-V.
Pfam; PF07654; C1-set; 3.
SMART; SM00409; IG; 2.
SMART; SM00409; IGc1; 3.
SMART; SM00407; IGc1; 3.
SMART; SM00407; IGc1; 3.
SMART; SM00406; IGV; I.
PROSITE; P850835; IG-LIKE; 4.
PROSITE; P8508290; IG-MHC; UNKONOM_2.
            PIR; A91668; GIHUNI.

HSSP; P01772; 2FB4.

GO; GO:0005576; C:extracellular; NAS.

GO; GO:0003823; F:antigen binding; NAS.

GO; GO:0006955; P:immune response; NAS.

InterPro; IPR007110; Ig-like.

InterPro; IPR003596; Ig_v.
                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE-77070367; PubMed=1002129;
Preker I., Schwarz J., Reichel W., Hilschmann N.;
"Rule of antibody structure. The primary structure of a monoclonal IgG1 immunoglobulin (myeloma protein Nie), I: purification and characterization of the protein, the L- and H-chains, the cyanogen bromide cleavage products, and the disulfide bridges.";
Hoppe-Seyler's Z. Physiol. Chem. 357:1515-1540(1976).
Hoppe-Seyler's Z. Physiol. Chem. 357:1515-1540(1976).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ponstingl H., Hilschmann N.;
"The rule of antibody structure. The primary structure of a monoclonal IgG1 immunoglobulin (mycloma protein Nie). III. The chymotryptic peptides of the H-chain, alignment of the tryptic peptides and discussion of the complete structure.";
hoppe-Seyler's Z. Physiol. Chem. 357:1571-1604(1976).
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                 DISULFIDE BOND.
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Pred. No. 1.1e
3; Mismatches
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1.1e-41;
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                                                                                               1 QVQLVESGGGVVQPGRSLRLSCAASGPTFSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY
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Search completed: ( Job time : 120.489

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                GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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| 78.1               |        |                    |                    |        | 78.9   | 78.9   | 78.9               | 79.3   | 79.3   | 79.6   | 79.6   | 79.7   | 79.7               | 79.8               |
| 120                | 138    | 121                | 98                 | 147    | 119    | 113    | 122                | 121    | 119    | 122    | 122    | 120    | 119                | 117                |
| 2                  | N      | ۲                  | N                  | Ŋ      | N      | N      | N                  | N      | N      | N      | _      | N      | N                  | N                  |
| S44111             | B56701 | CIHUHL             | PL0116             | I37780 | C36005 | S38490 | S31119             | I55673 | S31108 | 869910 | M3HUAM | S36278 | S31107             | S78486             |
| Ig heavy chain V-D |        | Ig heavy chain V-I | Ig heavy chain V-I |        |        |        | Ig heavy chain - h |        | heavy  |        | -      |        | Ig heavy chain - h | Ig heavy chain V r |

#### ALIGNMENTS

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Ig heavy chain V region - human (fragment)
C;Species: Homo sapiens (man)
C;Species: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
C;Accession: S31679
R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C.
submitted to the EMBL Data Library, June 1992
A;Description: Mechanisms that generate human immunoglobulin diversity operate from A;Reference number: S31585
A;Accession: S31679
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-134 <CUI>A;Residues: 1-134 <CUI>A;Cross-references: EMBL: Z14203; NID:g30965; PIDN:CAA78572.1; PID:g30966
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Status: preliminary; nucleic acid sequence not shown; translation A;Molecule type: mRNA A;Residues: 1-122 <RAA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bur. J. Immunol. 22, 247-251, 1992
A;Title: Restricted utilization of germ-line V(H)3 genes and short diverse third complem A;Reference number: S31104; MUID:92111633; PMID:1730252
A;Accession: S31117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ig heavy chain - human

C;Species: Homo sapiens (man)

C;Date: 02-Dec-1993 #sequence_revision 26-May-1995 #text_change 17-Mar-1999

C;Accession: S31117
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F;15-98/Domain: immunoglobulin homology <II</pre>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Raaphorst, F.M.; Timmers, B.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurmar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SS 119
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88.3%;
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86.1%;
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Pred. No. 5.9e-43;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 533.5;
Pred. No. 6e
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Length 134;
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Query Match

Score 533;

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C; Superfamily: ______C; Reywords: heterotetramer; ______CC; Reywords: heterotetramer; ______CC; Reywords: immunoglobulin homology F; 15-98/Domain: immunoglobulin homology
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A;Residues: 1-119 <SCH>
A;Cross-references: UNIPROT:QBWUK1; GB:M34026
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rischroeder Jr., H.W.; Wang, J.Y.
Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990
A;Title: Preferential utilization of conserved immunoglobulin A;Reference number: A36005; MUID:90349571; PMID:2117273
A;Accession: P36005
A;Status: preliminary
A;Status: preliminary
                                                                                                                                                A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-132 <CUI>
A;Crose-references: EMBL:Z14168; NID:g30999; PIDN:CAA78537.1;
A;Crose-references: EMBL:Z14168; NID:g30999; PIDN:CAA78537.1;
C;Superfamily: immunoglobulin V region; immunoglobulin homolog
C;Keywords: heterotetramer; immunoglobulin
F;30-113/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                            C;Accession: S31603

R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, submitted to the EMBL Data Library, June 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Cross-references: GDB:118731;
A; Map position: 14q32.33-14q32.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ig heavy chain V region (M49) - C; Species: Homo sapiens (man)
                                                                                                                                                                                                                                                                                               A; Reference number: S31585
A; Accession: S31603
                                                                                                                                                                                                                                                                                                                                                                                                                            Ig heavy chain V region - human
C;Species: Homo sapiens (man)
C;Date: 03-Mar-1994 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Map position: 14q32.33-14q32.33
C; Superfamily: immunoglobulin V region; immunoglobulin homology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Gene: GDB: IGH@; IGHDY1
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C;Accession: F36005
                                                                                                                                                                                                                                                                                                                                             A; Description: Mechanisms that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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Best Local
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Best Local :
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QVQLVESGGGVVQPGRSLRLSCAASGFTESSSGMHWVRQAPGKGLEWVAXISYDGSRKYY
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ilarity 86.6%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 532; DB 2
Pred. No. 8e-43;
2; Mismatches
                                                                   Score 531; DB 2;
Pred. No. 1.1e-42;
1; Mismatches 11
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A;Map position: 14932.33-14932.33
C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;15-98/Domain: immunoglobulin homology <IVM>
                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: mRNA
A; Residues: 1-121 < SCH>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Schroeder Jr., H.W.; Wang, J.Y.
Proc. Natl. Acad. Sci. U.S.A. 87, 6146-6150, 1990
A;Tille: Preferential utilization of conserved immunoglobulin heavy
A;Reference number: A36005; MUID:90349571; PMID:2117273
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A;Title: IgM kappa/landbda EBV human B cell clone: an early step of differentiation A;Reference number: S70442; MUID:93024508; PMID:1383695
A;Accession: S70442
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C;Species: Homo sapiens (man)
C:Date: 21-Dec-1990 #sequence_revision 21-Dec-1990 #text_change 09-Jul-2004
C;Accession: G36005
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
F;34-117/Domain: immunoglobulin homology <IMM>
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Ig heavy chain precursor V region (mu) - human (fragment)
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Best Local
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05; Conservative
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                         ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAKGVT--GSPTLDYWGQGTLVTVS
                                                                                                     QVQLVESGGGVVQPGRSLRLSCAASGFTFSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAKG-VTGSPTLDYWGQGTLVTVSS 119
  ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDRKDWGWALFDYWGQGTLVTVS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDHIVGATYFDYWGQGTLVTVSS
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85.8%; Pred. No. 1.3e-42;
ative 4; Mismatches 12;
                                                                                                                                                       86.0%; Score 529; DB 2; I
86.8%; Pred. No. 1.6e-42;
tive 1; Mismatches 13;
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R;Cuisinier,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ig heavy chain V region - human
C;Species: Homo sapiens (man)
C;Date: 03-Mar_1994 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R;Raaphorst, F.M.; Timmers, E.; Kenter, M.J.H.; van Tol, M.J.D.; Vossen, J.M.; Schuurmai Bur. J. Immunol. 22, 247-251, 1992
A;Title: Restricted utilization of germ-line V(H)3 genes and short diverse third complei A,Reference number: S31104; MUID:92111633; PMID:1730252
A,Accession: S31116
A,Status: preliminary; nucleic acid sequence not shown; translation not shown A;Molecule type: mRNA
A,Residues: 1-118 <RAA>
RESULT 11
$31598
Ig heavy chain V region -
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A;Note: the nucleotide sequence was submitted to the EMBL Data Library, October 1991
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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C;Date: 02-Dec-1993 #sequence
C;Accession: S31116
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Matches 104;
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                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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MBL Data Library, June 1992
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                                                                                                                                                                                                                                                                                                                                                  85.7%;
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Pred. No. 2.6e-42;
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human

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a, curbinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.; Tonnelle, C. submitted to the EMBL Data Library, June 1992
A;Description: Mechanisms that generate human immunoglobulin diversity operate from A;Reference number: S31585
A;Status. 531598
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Accession: PH1643
A;Molecule type: mRNA
A;Residues: 1-111 <HIL>
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A;Title: The structural basis of germline-encoded VH3 immunoglobulin binding to A;Reference number: PH1642; MUID:93301610; PMID:8315388
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C;Species: Homo sapiens (man)
C;Date: 24-Feb-1994 #sequence_revision 24-Feb-1994 #text_change 09-Jul-2004
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A; Residues: 1-135 < CUI>
A; Cross-references: EMBL: Z14170; NID: g31001; PIDN: CAA78539.1; PID: g31002
C; Superfamily: immunoglobulin V region; immunoglobulin homology
C; Keywords: heterotetramer; immunoglobulin
A;Status: preliminary A;Molecule type: DNA
                                   C;Date: 27-Jan-1995 #sequence_revision 27-Jan-1995 #text_chan c;Accession: 846390
C;Accession: 846390
R;Figini, M; Marks, J.D.; Winter, G.; Griffiths, A.D. J. Mol. Biol. 239, 68-78, 1994
A;Title: In vitro assembly of repertoires of antibody chains A;Reference number: 846390; MUID:94254092; PMID:8196048
A;Accession: 846390
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C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
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                                                                                                                                        ;Species: Homo sapiens (man)
;Date: 27-Jan-1995 #sequence_revision
;Accession: S46390
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Best Local
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91.0%;
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Pred. No. 2.7e-42;
0; Mismatches 15
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Pred. No. 4.2e-42;
                                                                                                                                                             27-Jan-1995 #text_change
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A;Residues: 1-114 <FIG>A;Residues: 1-114 <FIG>A;Ross-references: EMBL:231686; NID:g509782; PIDN:CAA83491.1; PID:g1335143 C;Superfamily: immunoglobulin V region; immunoglobulin homology C;Keywords: heterotetramer; immunoglobulin F;15-98/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                  Ig heavy chain V region - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 23-Jul-1999
C;Accession: S31674
R;Cuisinier, A.M.; Gauthier, L.; Boubli, L.; Fougereau, M.: Tonnelle, C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 15
S31674
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Cross-references: GB.S64471; NID:g236904; PIDN:AAB20011.1; PID:g236905
A;Experimental source: X-linked agammaglobulinemia patients, B lymphoblastoid
A;Note: sequence extracted from NCBI backbone (NCBIN:64471, NCBIP:64470)
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
F;15-98/Domain: immunoglobulin homology <IMM>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C;Accession: A49028
R;Timmers, E.; Kenter, M.; Thompson, A.; Kraakman, M.E.; Berman, Bur. J. Immunol. 21, 2355-2363, 1991
Bur. J. Immunol. 21, 2355-2363, 1991
A;Title: Diversity of immunoglobulin heavy chain gene segment revalues and the segment 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ig heavy chain V-III region - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 21-Jan-1994 #sequence revision 18-Nov-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     밁
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A;Cross-references: EMBL:Z14204; NID:g30967; PIDN:CAA78573.1; PID:g30968 C;Superfamily: immunoglobulin V region; immunoglobulin homology
                                                                      A; Molecule type: mRNA
A; Residues: 1-139 < CUI>
                                                                                                                                                                           A; Reference number: S31585
A; Accession: S31674
                                                                                                                                                                                                                                                                                        submitted to the EMBL Data Library, June 1992
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A; Residues: 1-133 <TIM>
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Pred. No. 5.
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| Search completed: October 13, 2005, 03:11:46<br>Job time : 25.515 secs | Qy 61 ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAK-GVTGSPTLDYWGQGTLVTVSS 119 | QY 1 QVQLVESGGGVVQPGRSLRLSCAASGFTFSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY 60 | Query Match 84.5%; Score 519.5; DB 2; Length 139;<br>Best Local Similarity 87.5%; Pred. No. 1.4e-41;<br>Matches 105; Conservative 1; Mismatches 13; Indels 1; Gaps 1; | C;Keywords: heterotetramer; immunoglobulin<br>F;34-117/Domain: immunoglobulin homology <imm></imm> |
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OM protein - protein search, using sw model
                                                                                            GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.
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Run on: October 13, 2005, 02:44:40;

Title: Perfect score: US-10-010-729A-7 615 ; Search time 122.575 Seconds (without alignments) 375.480 Million cell updates/sec

Scoring table: Sequence: BLOSUM62 1 QVQLVESGGGVVQPGRSLRL.....VTGSPTLDYWGQGTLVTVSS 119

Searched: 2105692 seqs, 386760381 residues Gapop 10.0 , Gapext 0.5

Total number of hits satisfying chosen parameters:

2105692

Y., .

Minimum DB Maximum DB seq length: 0 seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database : A\_Geneseq\_16Dec04:\*
1: geneseqp1980s:\*
2: geneseqp1990s:\*
3: geneseqp2000s:\*
4: geneseqp2001s:\*
5: geneseqp2001s:\*
6: geneseqp2003as:\*
7: geneseqp2003bs:\*
8: geneseqp2004s:\* geneseqp2001s:\*
geneseqp2002s:\*
geneseqp2003as:\*
geneseqp2003bs:\*
geneseqp2004s:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

| 25                | 2 4      | 23                 | 22                 | 21                 | 20                 | 19                 | 18                 | 17                 | 16       | 15                 | 14                | 13       | 12         | 11       | 10                 | 9        | 8                  | 7        | 6        | υ        | 4                  | ω                  | N                  | _        | Result<br>No.         |
|-------------------|----------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|----------|--------------------|-------------------|----------|------------|----------|--------------------|----------|--------------------|----------|----------|----------|--------------------|--------------------|--------------------|----------|-----------------------|
| 534               | 534      | 534                | 534.5              | 535                | 535                | 535                | 535.5              | 537                | 537      | 537.5              | 538               | 538      | 538.5      | 538.5    | 539                | 540      | 540.5              | 540.5    | 542.5    | 542.5    | 544                | 611                | 611                | 611      | Score                 |
| 86.8              | 86.8     | 86.8               | 86.9               | 87.0               | 87.0               | 87.0               | 87.1               | 87.3               | 87.3     | 87.4               | 87.5              | 87.5     | 87.6       | 87.6     | 87.6               | 87.8     | 87.9               | 87.9     | 88.2     | 88.2     | 88.5               | 99.3               | 99.3               | 99.3     | Query<br>Match Length |
| 119               | 117      | 117                | 118                | 519                | 244                | 117                | 223                | 121                | 117      | 583                | 119               | 117      | 241        | 118      | 119                | 121      | 121                | 120      | 490      | 254      | 123                | 119                | 119                | 119      |                       |
| 7                 | œ        | œ                  | σ                  | ហ                  | 7                  | 8                  | N                  | ω                  | ω        | œ                  | 7                 | æ        | 7          | თ        | σ                  | œ        | œ                  | σ        | w        | w        | 0                  | ω                  | ຫ                  | u        | BB                    |
| ADL91318          | ADO36345 | ADI22095           | AAE32097           | AAU81993           | ADG30422           | ADI22096           | AAY08598           | ADP22186           | ADI22094 | ABM82698           | ADL91327          | ADO36354 | ADG3 04 67 | ABG30463 | AAE32095           | ADP22284 | ADP47108           | ADA89174 | AAY56637 | AAY56287 | ABR55789           | ADI26654           | ABB07186           | ABB07169 | ID                    |
| Ad191318 VH chain |          | Adi22095 Anti-plat | Aae32097 Human VEG | Aau81993 Human sec | Adg30422 Human GMB | Adi22096 Anti-plat | Aay08598 Anti-huma | Adp22186 Human ant |          | Abm82698 Human dia | Adl91327 VH chain | Intra    | 'Human     | Human    | Aae32095 Human VEG | Human    | Adp47108 Human pho | Human    | hCAT1    |          | Abr55789 Heavy cha | Adi26654 Human ant | Abb07186 sHigM22 h | 9 sHig   | Description           |

| •         |           |           | ×         |           |           |           |           |           |           |           |           |           | •         |           |           |           |           | i         |              |
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| 85.9      | 85.9      | 86.0      | 86.1      | 86.1      | 86.1      | 86.2      | 86.2      | 86.2      | 86.2      | 86.2      | 86.3      | 86.3      | 86.3      | 86.3      | 86.4      | 86.4      | 86.5      | 86.6      | 00.0         |
| 249       | 249       | 115       | 252       | 252       | 121       | 472       | 124       | 123       | 123       | 123       | 252       | 252       | 126       | 119       | 135       | 116       | 119       | 223       | 177          |
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| Adg95328  | Abp44501  | Aab39519  |           |           | Adp47221  |           | Adp46956  | Adr72698  | Adr72704  | Adr72700  |           |           | -         | •,        | •••       | Adc60976  | Aau02515  | Ad170773  | 1221 Ednw    |
| Single ch | Human BLy | Anti-IL12 | Single ch | Human BLy | Human pho | Human mon | Murine he | Human mon | Human mon | Human mon | Single ch | Human BLy | Murine-ex | Human foe | Human ant | Human ant | Anti-adip | Anti-TNFa | מתוומנו ביוט |

### ALIGNMENTS

ABB07169 standard; protein; 119

ABB07169;

13-MAR-2002 (first entry)

sHigM22 heavy chain variable region clone A sequence.

Neuromodulatory; central nervous system; CNS; sHIgM22; LYM 22; AKJR4; ebvHigM MsI19D10; ebv HIgM CB2bG8; CB2iB12; CB2iE7; MSI19E5; virucide; antiparkinsonian; neuroprotective; nootropic; vulnerary.

Homo sapiens.

2.

WO200185797-A1

15-NOV-2001.

30-MAY-2000; 2000WO-US014902.

10-MAY-2000; 2000US-00568351.

(MAYO-) MAYO FOUND MEDICAL EDUCATION RES.

Rodriguez M, Miller DJ, Pease LR;

WPI; 2002-066596/09. N-PSDB; ABA94216.

Novel neuromodulatory agent (a human IgM monoclonal antibody), promoting neurite outgrowth, regeneration, remyelination and neuroprotection in central nervous system, useful to treat post-infectious encephalomyelitis.

Claim 23; Fig 17; 219pp; English.

RESULT 1
ABB07169
ID ABB0717
XX ABB071
XX Homo
XX Neuron
PN W0200
XX Homo
XX Homo
PN W0200
XX Homo
PN WPI7
PN Novel
PT neuri
PT encet
XX WPI7
PT neuri
PT encet
XX Clain
XX The invention provides a neuromodulatory agent (I) capable of promoting neurite outgrowth, regeneration, remyelination and neuroprotection in central nervous system (CNS). (I) is capable of inducing remyelination, promoting cellular proliferation of glial cells, and promoting Ca2+ signaling with oligodendrocytes. An humanised antibody to (I) can be selected from antibody sHIgM22 (LYM 22), ebvHigM MSI19D10, ebv HIgM CB2bB6, AKUR4, CB2iE12, CB2iE7 or MSI19E5. (I) is useful for stimulating remyelination of CNS axons, stimulating proliferation of glial cells in CNS axons, or treating demyelinating disease of CNS in a mammal in need of such therapy. (I) is capable of binding to structures and cells within

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CC CNS. (I) is preferably useful for treating a demyelinating disease of CNS CC of a mouse infected with Strain DA of Theiler's murine encephalomyelitis CC (TMEW) or for treating a human being having multiple sclerosis, or a CC human or domestic animal with a viral demyelinating disease, or a post-cural disease of CNS. (I) is also useful for an in vitro method of CNS stimulating the proliferation of glial cells from mixed cell culture. (I) CC is also useful for stimulating remyelination of CNS axons. The antibodies CC are useful for preventing infection by a bacterium, virus or like CC are useful for envelination or other neurodegenerative condition in a subject. Methods where (I) is administered to a patient are useful CC for treating multiple sclerosis, Parkinson's disease, Alzheimer's CC disease, Amyotrophic lateral sclerosis (ALS), a viral demyelinating CC disease, and other conditions in the CNS where nerves are CC damaged as by trauma. The present sequence represents the sHigM22 heavy CC chain variable region clone A amino acid sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200185797-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Neuromodulatory; central nervous system; CNS; sHIgM22; LYM 22; AKJR4; ebvHigM MsI19D10; ebv HIgM CB2bG8; CB2iE12; CB2iE7; MSI19E5; virucide; antiparkinsonian; neuroprotective; nootropic; vulnerary.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-MAR-2002
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The invention provides a neuromodulatory agent (I) capable of promoting neurite outgrowth, regeneration, remyelination and neuroprotection in central nervous system (CMS). (I) is capable of inducing remyelination, promoting cellular proliferation of glial cells, and promoting Ca2+
                                                                                                                                        Novel neuromodulatory agent (a human IgM monoclonal antibody), prorneurite outgrowth, regeneration, remyelination and neuroprotection central nervous system, useful to treat post-infectious
                                                                                                                                                                                                                                                                   Rodriguez
                                                                                                                                                                                                                                                                                                                                           10-MAY-2000;
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                                                                                                                                                                                                                   N-PSDB;
                                                                                                                          encephalomyelitis.
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                                                                                                                                                                                                                                                                    Miller DJ,
                                                                                                                                                                                                                                                                                                                                           2000US-00568351.
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                                                                                        17;
                                                                                     219pp;
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                                                                                      English
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Pred. No. 2.7e-49;
D; Mismatches 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           region clone B sequence.
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Matches 117; Conserv
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                                                                                                                                            07-JAN-1997;
28-MAY-1999;
30-MAY-2000;
                                                                                                                                                                                                     29-APR-1994;
08-AUG-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human; antibody; IgM; remyelination; neuronal growth; autoantibody; demyelination disease; multiple sclerosis; central nervous system;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     spinal cord injury.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        axon; glial cell proliferation;
Theiler's murine encephalomyelitis virus infection; CNS injury;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human anti IgM antibody sHIgM22 VH protein.
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                                          Rodriguez M,
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                                                                                  FOUND.
                                                                                                                         99US-00322862.
2000US-00580787.
2000US-00730473.
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                                            Miller
                                                                                                                                                                                    94US-00236520.
96US-00692084.
97US-00779784.
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                                            Pease LR;
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Pred. No. 2.7e-49;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
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WPI; 2004-119219/12

New human immunoglobulin M antibody for treating or preventing demyelinating disease of the central nervous system in a human domestic animal, such as multiple sclerosis.

Claim 7; Fig 35; 159pp; English.

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RESULT 4
ABR55789
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Cimmunocompetent host with an antibody (I) produced by injecting an antibody, where the peptide comprises a human anti-Igw antibody fragment (I) given in the specification, or active fragments. Also included are stimulating remyelination of central nervous system (CNS) axons in a mammal (comprising administering a monoclonal antibody, or mixtures, color, active fragments, or recombinant antibodies derived from it, characterised by their ability to bind structures and cells within the CNS, including oligodendrocytes), stimulating the proliferation of glial cells in CNS axons in a mammal (comprising administering a monoclonal antibody, or mixtures, monomers, active fragments, or recombinant can tibodies derived from it, characterised by their ability to bind structures and cells within the CNS), treating or preventing a functional antibody, or mixtures, monomers, active fragments, or the bind structures and cells within the CNS), treating or preventing a compocional antibody, or mixtures, monomers, active fragments, or recombinant antibodies derived from it, characterised by their ability to faxons of the CNS), stimulating, in vitro, the proliferation of GNS axons in a mammal (comprising administering a compocional antibody, or mixtures, monomers, active fragments of glial coll axons of the CNS), stimulating, in vitro, the proliferation of GNS axons in a mammal (comprising administering coll antibody), a peptide analogue, hapten, or active fragment of it, where the DNA sequence consists of a sequence encoding an anti-IgM antibody), a probe capable of screening for the antibody, an assay for screening drugs and other agents for the ability to modulate the production or mimic the cNS comprising drugs and to stimulate remyelination of CNS axons, and to stimulate soft them, a cativities of mass apprision of the CNS using the antibody mucleic acids or recombinant virus transformed with recombinant antibody mucleic acids or combinant of the CNS axons, and to stimulate the production or mimic the color and to treat a sp
                                                                                                                                                                                                                                                                                                                                                              Matches 118; Conservative
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Best Local
                 02-SEP-2003
                                                                                           ABR55789 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 119
                                                                                                                                                                                                 61
                                                                                                                                                                                                                    61 ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAKGVTGSPTLDYWGQGTLVTVSS 119
                                                                                                                                                                                                                                                                                                                      μ
                                                                                                                                                                                                                                                                                                                                                          h 99.3%; Score 611; DB 8; Similarity 99.2%; Pred. No. 2.7e-49; 18; Conservative 0; Mismatches 1
                                                                                                                                                                                                                                                                                                 QVQLVESGGGVVQPGRSLRLSCAASGFTFSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY 60
                                                                                                                                                                                                                                                                           QVQLVESGGGVVQPGRSLRLSCAASGFTFSSSGMHWVRQAPGKGLEWVAVISYDGSRKYY 60
                                                                                                                                                                                           ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAKGVTGSPTLDYWGQGTLVTVSS 119
            (first entry)
                                                                                           123
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RESULT 5
AAY56287
ID AAY5

AAY56287 standard; protein; 254

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Best Local Simi
Matches 107;
                                    The invention relates to a specific binding agent, which comprises at least one peptide selected from any of 62 peptides (ABR55769-830) or its fragment. The binding agents are antibodies that recognize and bind to angiopoletin-2 (Ang-2). The specific binding agent, particularly the antibody, is useful for inhibiting undesired angiogenesis, treating cancers, inhibiting undesired angiogenesis, modulating or inhibiting Ang-2 activity, modulating vascular permeability or plasma leakage, or treating a disease (e.g. ocular neovascular disease, obesity, haemangioblastoma, haemangioma, arteriosclerosis, inflammatory disease, inflammatory disease, atherosclerosis, endometriosis, neoplastic disease, bone-related disease, or psoriasis) in a mammal. The present sequence represents a heavy chain variable region of an anti-Ang-2
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                                                                                                                                                                                                                                                                                Sequence 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New specific binding agents (i.e. anti-Angiopoietin-2 antibodies), useful for inhibiting undesired angiogenesis, or treating e.g. cancers, obesity, hemangioma, arteriosclerosis, atherosclerosis or endometriosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         11-OCT-2001; 2001US-0328604P
10-OCT-2002; 2002US-00269805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; Page 91; 161pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                11-OCT-2002; 2002WO-US032613
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gynaecological; antiinflammatory; osteopathic; antipsoriatic; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Heavy chain variable region of anti-Ang-2 antibody
                                                                                                                                                                                                          y Match 88.5%;
Local Similarity 87.0%;
nes 107; Conservative
                     117 VSS 119
                                                                                                                                                           1 QVQLVESGGGVVQPGRSLRLSCAASGFTFSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY
                                                                                         61 ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAKGVT----GSPTLDYWGQGTLVT 116
                                                                 61 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAKGPVDFDYGDYAIDYWGQGTLVT
                                                                                                                                       QVQLQESGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVAVISYDGSNKYY
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26. .36
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                                                                                                                                                                                                          Score 544; DB 6; Length 123; Pred. No. 5e-43; 1; Mismatches 11; Indels
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RESULT 6
AAY56637
ID AAY5
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AC AAX5
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AC AAX5
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AC AAX5
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Best Local
                                                                                                                                                                                                                                                                                                                             Matches 105;
                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention describes a virus-like particle or gene delivery vehicle comprising a ligand capable of binding to a human amino acid transporter. The method is useful for the target delivery of substances to cells e.g. gene therapy. A human cationic amino acid transporter (hCAT1) targeted adenovirus is useful for local applications of adenoviral vector e.g. in patients with restenosis following balloon angioplasty where smooth muscle cells need to be transduced with an adenoviral vector carrying the CeNOS cDNA. More efficient transduction of tissues can be carried out therefore resulting in lower multiplicity's of infections that can be used and therefore less vector associated toxicity to the tissues surrounding the target cells. AAZ38737 to AAZ38770, and AAY56264 to AAY56287 represent sequences used in the exemplification of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAY56287;
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 254 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 13; Fig
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New virus-like particle or gene delivery vehicle, useful for gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-001283/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-MAY-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-MAY-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     smooth muscle cell; transduction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human cationic amino acid transporter; hCAT1; targeted delivery; gene delivery; virus-like particle; retroviral envelope molecule;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HCAT1 clone 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              08-FEB-2000
Virus-like particle; ligand; gene delivery; envelope protein; hCAT1;
                              hCAT1 binding human antibody molecule protein sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (INTR-) INTROGENE BV
                                                              22-FEB-2000
                                                                                                                          AAY56637 standard; protein; 490
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .nfection;
                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                       8
                                                                                                                                                                                                                                   61
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAZ38770.
                                                                                                                                                                                                                                   ADSVKGRFTISRDNSKNTLYLOMNSLTAXDTAVYYCAKGVTGSPT-LDYWGQGTLVTVSS 119
                                                                                                                                                                                                                                                                                  QVQLVESGGGVVQPGRSLRLSCAASGFTFSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY
                                                                                                                                                                                                                                                                 QVQLVQSGGGVVQPGRSLRLSCAASGFTFSSYAMHWVRQAPGKGLEWVAVISYDGSNKYY
                                                                                                                                                                                                       ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGITVTKSRFDYMGQGTLVTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gene therapy; restenosis; balloon angioplasty;
                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                         invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16; 66pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antibody variable heavy chain protein sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98EP-00201693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98EP-00201693.
                                                                                                                                                                                                                                                                                                                                            88.2%;
                                                                                                                                                                                                                                                                                                                               5
                                                                                                                                                                                                                                                                                                                                            Score 542.5;
Pred. No. 1.5
                                                                                                                            B
                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                               .5e-42;
                                                                                                                                                                                                                                                                                                                                                          DB 3;
                                                                                                                                                                                                                                                                                                                               9;
                                                                                                                                                                                                                                                                                                                                                            Length 254;
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                                                                                                                                                                                                                                                                                                                               Gaps
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RESULT 7 ADA89174

ADA89174 standard;

protein; 120

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XXXXX

20-NOV-2003 ADA89174;

(first entry)

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CC vehicle (I) provided with a ligand capable of binding to a human amino CC acid transporter. (I) is used to deliver genes to human cells or primate CC cells that express the hCAT1 amino acid transporter, such as endothelial. CC haematopoietic or smooth muscle cells, as part of a gene therapy regime. CC The vectors are especially useful for providing local applications of CC adenoviral vector to patients with restences following balloon CC angioplasty, where smooth muscle cells need to be transduced with ceNOS CC CDNA, for example. (I) may also be used to pseudotype recombinant type CC retrovirus including murine leukemia retroviruses and lentiviruses. In CC many cell types as possible is required. The new gene delivery vehicles transduce DNA more efficiently and specifically into tissues that are CC compared to a wildtype adenoviral vector. This increased specificity CC compared to a wildtype adenoviral vector. This increased specificity CC results in lower multiplicities of infection which can occur with prior CC art vectors, so preventing tissue toxicity. In addition the new vectors and tissues, which increases their efficiency for use as gene CC increases the potential host cells that are in low abundance in cell therapy vehicles. The alteration of the ligand on the viral envelope CC increases the potential host cell range of these vehicles. The present or present invention a hCAT1 binding human antibody molecule from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches 105;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                human cationic amino acid transporter 1; retrovirus; adenovirus; targeted delivery; gene therapy; balloon angioplasty.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 490 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           patients
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-025491/03.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
319
                                                                                                                                                                                                                             259
                                                                             61 ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAKGVTGSPT-LDYWGQGTLVTVSS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention
                                                                                                                                                                                                                                                                                               QVQLVBSGGGVVQPGRSLRLSCAASGFTESSSGMHWVRQAPGKGLEWVAXISYDGSRKYY
                                                                                                                                                                                                                             UNITED TO STANDARD ST
ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARGITVTKSRFDYWGQGTLVTVSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 542.5; DB 3
Pred. No. 3.1e-42;
5; Mismatches 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                             119
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                 378
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Human antibody 1A9 heavy chain amino acid sequence

SEQ ID NO:18

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                                                                                                                                                                                                                                                                                                                                                        CC (Ig) heavy chain variable (VH) domain and an Ig light chain variable (VL) CC domain. The protein binds a complex comprising a major histocompatibility CC complex (MHC) and a peptide, does not substantially bind the peptide complex (CC MUCL). Tak or hTERT. Also described: (1) a pharmaceutical composition of the MHC. The peptide is a peptide fragment of gp100, CC MUCL). Tak or hTERT. Also described: (1) a pharmaceutical composition of comprising the novel protein and a carrier; (2) a cytotoxic T cell comprising the novel protein and a carrier; (2) a cytotoxic T cell comprising the bound peptide, does not substantially bind the MHC in the absence of the MHC, and does not substantially bind the MHC comprising a first segment that encodes the Ig variable domain; (4) a comprising a first segment that encodes the Ig variable domain; (4) a comprising a first segment that encodes the Ig variable domain; (4) a comprising a first segment that encodes the Ig variable domain; (4) a comprising a first segment that encodes the Ig variable domain; (4) a convel protein; (5) a transgenic animal whose genome includes heterologous nucleic acid sequences that encode the protein; (6) identifying the covel protein; (8) a transgenic animal whose genome includes heterologous concleic acid sequences that encode the protein; (6) identifying the covel protein; (8) ablating or killing a target cell that complex in a sumple. A protein of the invention has cytostatic activity, and composition for treating or preventing a cancerous disorder. The present of composition for treating or preventing a cancerous disorder. The present of composition for treating or preventing a cancerous disorder. The present of composition for treating or preventing a cancerous disorder. The present of composition for treating or preventing a cancerous disorder. The present of composition for treating or preventing a cancerous disorder.
                                                                                                                                                                                  Matches
                                                                                                                                                                                                     Query Match
Best Local
                                                                                                                                                                                                                                                                         Sequence 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention describes a protein comprising an immunoglobulin (Ig) heavy chain variable (VH) domain and an Ig light chain variable ('
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Fig 3B; 224pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hoogenboom HRJM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-FEB-2002; 2002US-0358994P
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immunoglobulin; Ig; heavy chain variable domain; light chain variable domain; major histocompatibility complex; MHC; gp100; MUC1; TAX; hTERT; cytostatic; gene therapy; cancerous disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (DYAX-) DYAX CORP.
(TECR ) TECHNION RES & DEV FOUND LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein comprising an immunoglobulin heavy chain variable (VH) an immunoglobulin light chain variable (VL) domain, useful for paring a composition for treating or preventing a cancerous disc
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                                                                                                                                                                                  106; Conservative
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                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                            complex where the peptide component in as peptide fragment
                               ADSVKGRFTISRDNSKVTLYLQMNSLTAXDTAVYYCAKGVTGS-PTLDYWGQGTLVTVSS
                                                                                                                                  QVQLVESGGGVVQPGRSLRLSCAASGFTFSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY 60
ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDYYGDYALLDYWGQGTLVTVSS
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                                                                                                                                                                                                   87.9%;
                                                                                                                                                                           Score 540.5; DB 6;
Pred. No. 1e-42;
2; Mismatches 11;
                                                                                                                                                                                                                         Length 120;
                                                                                                                                                                                Indels
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09-SEP-2004 (first entry)

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RESULT 8 ADP47108
                                          RESULT 9'
                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
Matches ... 105; Conserv
                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                   The invention comprises a human monoclonal antibody that binds to phospholipase A2 (PLA2). The monoclonal antibody of the invention is useful in the preparation of a medicament for the treatment of inflammatory and degenerative disorders stemming from inflammatory reactions in the joints, skin, and blood vessels, arthritis, psoriasis, asthma, Alzheimer's disease, atheroscierosis, and restenosis. The present amino acid sequence represents the heavy chain from a monoclonal antibody that is specific for the human phospholipase A2 (PLA2) enzyme.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ABGE-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New human monoclonal antibody that binds to phospholipase A2 (PLA2), useful for treating inflammatory conditions, e.g. arthritis, psorias asthma, Alzheimer's disease, atherosclerosis, or restenosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   joint inflammatory reaction; skin inflammatory reaction; blood vessels inflammatory reaction; arthritis; psoriasis; asthma; Alzheimer's disease; atherosclerosis; restenosis; heavy chain.
ADP22284;
                              ADP22284 standard;
                                                                                                                                                                                                                                                                                                        Sequence 121 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Landes GM, Haak-Frendscho M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    02-DEC-2003; 2003WO-US038234.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    human; monoclonal antibody; phospholipase A2; PLA2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human phospholipase A2-specific monoclonal antibody heavy chain #11.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   inflammatory disorder; degenerative disorder;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nocerini MR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABGENIX INC.
LEXICON GENETICS
                                                                                                                                                                                                 QVQLVESGGGVVQPGRSLRLSCAASGFTFSSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY
                                                                                                                                ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAKGVTGSP-TLDYWGQGTLVTVSS
                                                                                                                                                                           QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVAAIWYDGSNKWY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQ ID NO 23; 128pp; English.
                                                                                                              ADSVKGRFT1SRDNSKNTLYLQMNSLRAEDTAVYYCARGGTGTPGAFD1WGQGTMVTVSS
                                                                                                                                                                                                                                        87.9%;
ilarity 87.5%;
Conservative
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                            protein;
                              121
                                                                                                                                                                                                                                        Score 540.5; DB 8;
Pred. No. 1e-42;
4; Mismatches 10;
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                                                                                                                                                                                                                                                                    Length
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                                                                                                                                                                                                                                                                         121;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       psoriasis,
                                                                                                                                                                                                                                         Gaps
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Human anti-TNFa antibody heavy chain variable region

SEQ

ID NO:190.

HANNARAN MANARAN KANARAN KANAR eating-disorder; immunomodulator; immunosuppressive; nephrotropic; neuroprotective; vasorropic; antiapoptotic; TNFa antagonist; TNF induced apoptosis; neoplastic disease; breast cancer; ovarian cancer; bladder cancer; lung cancer; glioblastoma; stomach cancer; endometrial cancer; kidney cancer; colon cancer; pancreatic cancer; prostrate cancer; immuno-mediated inflammatory disease; restenosis; autoimmune disease; Crohn's disease; graft-host reaction; septic shock; cachexia; anorexia; multiple sclerosis. anti-TNFa antibody; anabolic; antiarteriosclerotic; antiartaria antibacterial; antiinflammatory; antipsoriatic; antirheumatory human; monoclonal antibody; tumour necrosis factor-alpha; TNFa; antiarteriosclerotic; antiarthritic;

17-JUN-2004 WO2004050683-A2

02-DEC-2003; 2003WO-US038281.

02-DEC-2002; 2002US-0430729P

ABGENIX INC

Haak-Frendscho M, Babcook JS, Kang JS, Faggioni R, JS, Foord O, Green L, Feng X, Klakamp Rathanaswami P, Pigott C, Liang ML, Le Paggioni R, Senaldi G, Qlaojuan JS; Ø

N-PSDB;

New recombinant human monoclonal antibody that specifically binds to Tumor Necrosis Factor-alpha, useful for treating neoplastic disease such as cancers, or immuno-mediated inflammatory diseases such as rheumatoid

Example 10; SEQ ID NO 190; 213pp; English

CC (a) a heavy chain complementarity determining region 1 (CDR1) having the CC two fully defined 5 amino acid sequence (S1, ADP22417) or (S2, ADP22421); CC and (b) a light chain CDR1 having the two fully defined 1 amino acid Sequence (S3, ADP22418) or (S4, ADP22424). Also described: (1) assaying (MM1) the level of TNFa in a patient sample, comprising contacting with CC (I), and detecting the level of binding between the antibody and TNFa in the sample; (2) a composition comprising the antibody or its functional CC fragment and a carrier; (3) treating (M2) an animal suffering from a CC enoplastic, or an immuno-mediated inflammatory disease by selecting an CC animal in need of treatment for the disease by administering the human CC monoclonal antibody of (I); and (4) inhibiting (M3) TNFa induced CC apoptosis in an animal by selecting an animal in need of treatment for CC (I). (I) has anabolic, antiarteriosclerotic, antiarthritic, eating-CC (I). (I) has anabolic, antiarteriosclerotic, pancreatic cancer, and prostrate cancer; or immuno-mediated inflammatory diseases such as rheumatoid arthritis, glomerullonephritis, atherosels psoriasis, restrates, autoimmune disease, Crohn's disease, graft-host reactions, septic shock, cachexia, anorexia, and multiple sclerosis. The present sequence represents a human anti-TNFa antibody heavy chain variable region, which is used in the The present invention describes a human monoclonal antibody (I) that specifically binds to tumour necrosis factor-alpha (TNFa) and compri(a) a heavy chain complementarity for a line of the comprise of the complementarity for the comprise of the comprise 얁 comprises:

밁 S 밁 Ş 밁 Matches Query Match Best Local Similarity AAE32095 standard; protein; 119 106; 119 121 S 121 61 5 Н \_ QVQLVESGGGVVQPGRSLRLSCAASGFTESSSGMHWVRQAPGKGLEWVAXISYDGSRKYY 60 ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAREGDYGGNPYFDYWGQGTLVTVS ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAK--GVTGSPTLDYWGQGTLVTVS 118 QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVAVIWYDGSNKYY 8 119 Conservative 87.8%; 2 Score 540; DB 8; Pred. No. 1.2e-42; Mismatches Length 121; Indels 2; Gaps 60

24-MAR-2003 (first entry

Human VEGF-2 hybridoma antibody #1

prollferative disorder; tumour; breast; cancer; brain; prostate; colon; lymphangioma; infection; Kaposi's sarcoma; psoriasis; immunosuppressive; rheumatoid arthritis; diabetic retinopathy; gene therapy; antimicrobial; cytostatic; ophthalmological; antibody; autoimmune disease. Human; vascular endothelial growth factor; VEGF-2; inflammatory disease;

Homo sapiens.

WO200283849-A2

24-OCT-2002

12-APR-2002; 2002WO-US011404

13-APR-2001; 2001US-0283391P. 07-SEP-2001; 2001US-0317600P.

(HUMA-) HUMAN GENOME SCI INC

Rosen CA, Albert VR, Ruben SM, Wager

N-PSDB; AAD49586. WPI; 2003-093007/08

New vascular endothelial growth factor (VEGF) - 2 antibodies, for treating, preventing or ameliorating a disease or disorder, such as inflammatory diseases, proliferative disorders, autoimmune disorders retinopathy. õ

Disclosure; Page 394; 399pp; English

RESULT 10
AAR32095
ID AAR322095
AC AAR32
XX AAR32
XX AAR32
XX AAR32
XX AAR32
XX Human
XX Human
XX Homo
XX Homo
XX WO200
XX W0200
XX I2-AI
XX I1-AI
XX I1-AI
XX II-AI
XX WPI
XX WP The invention relates to vascular endothelial growth factor (VEGF)-2 antibodies. VEGF-2 antibodies are useful for treating, preventing or ameliorating a disease or disorder, such as inflammatory diseases or disorders, proliferative disorders, tumours, tumour metastasis, breast cancer, brain cancer, prostate cancer, colon cancer, lymphangioma, an infectious disease, Kaposi's sarcoma, an autoimmune disease, rheumatoid arthritis, psoriasis, diabetic retinopathy, a disease or disorder associated with aberrant VEGF-2 (receptor) expression, or a disease or disorder associated with the lack of VEGF-2 (receptor) function. The antibody is also useful for detecting, diagnosing, prognosing, or monitoring cancers and other hyperproliferative disorders. VEGF-2 is also used antibody

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RESULT 11
ABG30463
ID BG30463
AC ABG30

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                                                     The invention relates to a human monoclonal antibody or fragment capable of specifically binding to a human CD40 antigen (a glycoprotein expressed on the surface of human B cells), where the antibody or fragment is free of significant agonistic activity when it binds to the CD40 antigen, and the growth or differentiation is inhibited. The fragments comprise the complementarity determining region (CDR) of the light and heavy chains of the monoclonal antibodies secreted by a hybridoma consisting of 1588, CC 20C4, 12D9, 9F7 and 13B4. Also included are the nucleic acids encoding the antibody (or fragments). The antibodies or fragments are used for inhibiting proliferation, growth or differentiation of a normal human B CC cells and to inhibit antibody production by B cells. They may also be cells and to inhibit antibody production by B cells. They may also be cerythematosus, psoriasis, multiple sclerosis, inflammatory bowel disease (Crohn's disease), rheumatoid arthritis, and lymphoma (especially Non-CC Hodgkin's Lymphoma). The present sequence represents the antibody 9F7 heavy chain variable region VH.1
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A human anti-CD40 monoclonal antibody or fragment useful for inhibiting proliferation, growth or differentiation of a normal human B cells and treating autoimmune disease such as rheumatoid arthritis or systemic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         02-OCT-2000; 2000US-0237556P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6; Fig 14; 75pp;
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Sequence 118 AA;

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RESULT 12
ADG30467
ADG30467
AC ADG3C
XX
AC
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Best Local Similarity

Matches 107; Conserv
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                                                                                                                                                                                                            Query Match 87.6%;
Best Local Similarity 88.2%;
Matches 105; Conservative
                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a novel antibody that specifically binds to a GMAD polypeptide comprising a first amino acid sequence that is at least 95% identical to a second amino acid sequence of a VH CDR (complementarity determining region) or VL CDR of an scPv (single chain antibody molecule). The antibody of the invention demonstrates antidiabetic activity and may be useful for diagnosing, monitoring, treating, preventing or ameliorating type II diabetes. The current sequence is that of the human scPv protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antibody that specifically binds to GMAD polypeptide, useful for diagnosing, monitoring, treating, preventing or ameliorating type II diabetes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-APR-2002; 2002US-0368813P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GMAD; VH; CDR; complementarity determining region; VL; single chain antibody; antidiabetic; type II diabetes;
                                                                                                                                                                                                                                                                                                                           Sequence 241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 2; SEQ ID NO 100; 410pp; English.
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61
                                                61 ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAKGVTGSPTLDYWGQGTLVTVSS 119
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  ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAKRAAAG-TLDYWGQGTTVTVSS
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                                                                                                                                                                                                               2;
                                                                                                                                                                                                               Score 538.5; DB 7;
Pred. No. 3.4e-42;
2; Mismatches 11;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 538.5; DB 5
Pred. No. 1.6e-42;
1; Mismatches 10
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                                                                                                                                                                                                                  Indels
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human;
                                                                                                                                                                                                               1;
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RESULT 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention relates to a novel method of determining the ability of an cimmunoglobulin single domain to bind to a target in an intracellular cenvironment comprising assessing the intracellular interaction between the immunoglobulin domain and the target by monitoring the signal. The method comprises providing a first molecule and a second molecule, where central interaction of the first molecule and a second molecule, where the stable interaction of a signal; providing a single intracellular immunoglobulin central intracellular immunoglobulin domain which is associated with the first molecule, where the single intracellular interaction of the signal, and assessing the intracellular interaction of the signal. The methods are useful for determining the ability of an complobulin single domain to bind to a target in an intracellular cenvironment. The present sequence is that of a single chain variable continual (scFv) protein which was used to illustrate the method of the
                                                                                                                                                                                                                                                         Query Match
Best Local Sim
Matches 107;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; SEQ ID NO 18; 66pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        monitoring the signal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Determining the ability of an immunoglobulin single domain to bind to target in an intracellular environment by assessing the intracellular interaction between the immunoglobulin domain and the target by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2004-431946/40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-NOV-2002; 2002GB-00026729.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   14-NOV-2003; 2003WO-GB004942.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-JUN-2004
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO2004046185-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               single chain variable fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26-AUG-2004
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                                                                                                                                                                                                                                                                                                                                                                                         Sequence 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             immunoglobulin single domain; intracellular environment;
intracellular interaction; immunoglobulin domain; scFv;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Intracellular interaction-related scFv protein SeqID18.
                                                                                                                                                                                                                                                                                              Local Similarity
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                                                                                                                                                              QVQLVESGGGVVQPGRSLRLSCAASGFTFSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY
                                                         ADSVKGRFTISRDNSKNTLYLOMNSLTAXDTAVYYCAKGVTGSPT-LDYWGQGTLVTVSS
   ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAK---ASPLHFDYWGQGTLVTVSS
                                                                                                                            QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGWHWVRQAPGKGLEWVSVISYDGSNKYY
                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                       87.5%;
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                                                                                                                                                                                                                                                            1;
                                                                                                                                                                                                                                                                                       Score 538; DB 8;
Pred. No. 1.7e-42;
                                                                                                                                                                                                                                                               Mismatches
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RESULT 14
ADL91327
ID ADL91327
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                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 107; Conserv
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15-NOV-2002;
15-NOV-2002;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Use of an intracellularly binding immunoglobulin comprising at least one antibody variable chain, in preparing a medicament for degrading one or more specific antigens, or for treating specific antigen positive cancer, e.g. leukemia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antibody; variable chain; cytostatic; cytoplasmic degradation; intracellular relocation; specific antigen positive cancer; lelymphoma; intracellularly binding immunoglobulin; BCR-ABL.
                                                                                                                                                                                                                                                                                                                                                                                               Sequence
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                                         ADSVKGRFTISRDNSKNTLYLQMNSLTAXDTAVYYCAKGVTGSPT-LDYWGQGTLVTVSS
                                                                                                                                                                 QVQLVESGGGVVQPGRSLRLSCAASGFTESSSGMHWVRQAPGKGLEWVAXISYDGSRKYY
ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAK---ASPLHFDYWGQGTLVTVSS
                                                                                                                              QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVSVISYDGSNKYY
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                                                                                                                                                                                                                                                                                               87.5%;
                                                                                                                                                                                                                                                                 Score 538; DB 7;
Pred. No. 1.8e-42;
1; Mismatches 8
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                                                                                                                                                                                                                                                                                                                                  Length 119;
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                                                                                                                                                                                                                                The invention relates to novel diagnostic and therapeutic polynucleotides CC selected from one of the 2722 sequences defined in the specification. A CC polynucleotide of the invention may have a use in gene therapy. The human CC diagnostic and therapeutic polynucleotides (dithp) or polypeptides may be cc used to diagnose a particular condition, disease or disorder associated CC with human molecules, e.g. cell proliferative disorder, endocrine CC autoimmune/inflammatory disorders, gastrointestinal disorder, endocrine CC disorder, neurological disorders, gastrointestinal disorder, or infections caused by virus, bacteria, fungi or parasite. The dithp CC molecules may also be used in genetic mapping, in identifying individuals CC from minute biological samples, in detecting single nucleotide polymorphisms, as molecular weight markers, and for somatic or germline CC gene therapy. The present sequence represents a dithp protein of the convention. Note: The sequence data for this patent is not represented in the printed specification, but was obtained in electronic format directly from WIPO at www.wipo.int/pct/en/sequences/listing.htm
                                                                                                                                                 Query Match
Best Local S
Matches 107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Schmidt JP, Wright RJ, Bruns CM, Marjanovic MM, Shen F; Harthshorne TA, Suchorolski MT, Altus CM, Pitts SJ, Elder LV; Mooney EM, Delegeane AM, Panesar IS, Banville SC, Reddy TP; Stevens KA, Blanchard JL, Panzer SR, Wang X, Au AP, Gerstin EH; Peralta CH, Anderson SB, Rioux P, Shen EJ, Wu MC, Stuve LL; Lagace RE, Spiro PA, Stewart EA, Wingrove J, Vitt UA, Kirton ES Xu Y, Kwong M, Policky JL, Hurwitz BL, Ma Y, Jackson JL, Gietzl Patury S, Shi X, Suarez CJ;
                                                                                                                                                                                                                                Sequence 583
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 27; Page; 190pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New diagnostic and therapeutic polynucleotides and polypeptides, useful in diagnosing a condition, disease or disorder associated with human molecules, e.g. autoimmune or inflammatory disorders, in gene therapy or in gene mapping.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2004-329368/30.
N-PSDB; ACN41350.
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12-SEP-2002; 2002US-0410260P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene therapy; human diagnostic and therapeutic polynucleotide; dithp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human diagnostic and therapeutic pprotein SEQ ID NO:2947.
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                                                                                                                                                 al Similarity 87.4%;
al Similarity 87.7%;
107; Conservative
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                     ADSVKGRFTISRDNSKNTLYLOMNSLTAXDTAVYYCAKGVTGSPT---LDYWGQGTLVTV 117
                                                                       QVQLVESGGGVVQPGRSLRLSCAASGFTFSSSGMHWVRQAPGKGLEWVAXISYDGSRKYY 60
ADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDT--PPTCDYFDYWGQGTLVTV
                                                                                                                                                                                                                                A,
                                                                                                                                               Score 537.5; DB 8;
Pred. No. 1.1e-41;
1; Mismatches 9;
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'', Stuve LL;
''A, Kirton ES;
''TA, T'., Gietzen D;
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Db 138 SS 139

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